GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen

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Minimum DB
Maximum DB
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence:
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Total number of hits satisfying chosen parameters:
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pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                  Score
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Match
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112.614 Million cell updates/sec
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Gapop 10.0 ,
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59
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1: pir1:*
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A58718
S78420
A60589
B46453
                                                                                                                                     PW0002
A56029
S27873
                           PT0238
SPPGNK
C61033
                                                                                                                  A38841
S35490
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                                                                                                                                                                                                                                                                                                                               SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1581
                                                                                                                                                                                              ribosomal protein
sperm-activating p
e antien p20e pre
alpha-myosin heavy
cytochrome b559 co
Ig heavy chain C r
coat protein beta
                  COI intron 16 prot peptida NADH2 dehydrogenas 1g heavy chain CRD neuromedin K pig ranatachykinin C tetracenomycin A2
                                                                                                        hypothetical prote
rhodopsin homolog
type II site-speci
translation elonga
                                                                                                                                                                                                                                                                    Ig heavy chain V r carnocin UI49 - Ca
                                                                                                                                                                             placental lactogen
pilE protein - Esc
                                                                                                                                                                                                                                                                                                  Description
                                                                                                glycoprotein H-a
                                                                                                                                                         polyphosphate-gluc
chlorophyll a/b-bi
                                                                                                                                                N-methylpurine DNA
                                                                                       -aminoimidazole r
proline-ric
                                                                                                                                                                                                                                                                                                                                                                                                          Ig heavy chain V region - human
(;Species: Homo sapiens (man)
(;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 16-Aug-1996
(;Accession: S21205
R;Makiya, R.; Stigbrand, T.
Bur. J. Biochem. 205, 341-345, 1992
Bur. J. Biochem. 205, 341-345, 1992
A;Title: Placental alkaline phosphatase has a binding site for the human immunoglobulin-large ference number: S21205; MUID:92209522; PMID:1555592
A;Accession: S21205
                                                                                                                                                                                                                                                                                                                                                                      A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-12 «MAK»
C;Keywords: heterotetramer; immunoglobulin
  RESULT
S78420
                                                                      Query Match
Best Local Similarity
**~*hes 4; Conserva
                                                                                                                  A;Molecule type: protein
A;Residues: 1-7 <5TO>
C;Keywords: antibiotic; lanthionine
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                                         SEIQ 5
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                                                                                                                                                                                                                                         Carnobacterium sp. (fragment)
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Ig H chain V-D-J r	PH1618	N	ထ		13
	JS0316	2	œ	22.0	13
al prot	T14906	~	00		13
d syntha	A20186	Ŋ	ወ	٠	13
RNA	T41235	2	12	23.7	14
۲,	PH1189	N	12	23.7	14
1 receptor a	PH1187	N	12	23.7	14
1 receptor	PH1190	N	12	23.7	14
yanın 1 - gr	C60529	N	12	23.7	14
potB protein - Sai	S71034	N	12	23.7	14
dad/	G49410	N	12	23.7	14
transcription fact	\$25485	2	12	23.7	14
	S41909	4.	11	23.7	14
T-cell receptor be	PH0924	Ŋ	11	23.7	14
T-cell receptor be	PH0904	2	11	23.7	14

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C;Accession: A58718
R;Stoffels, G.; Nissen-Meyer, J.; Gudmundsdottir, A.; Sletten, K.; Holo, H.; Nes, Appl. Environ. Microbiol. 58, 1417-1422, 1992
A;Title: Purification and characterization of a new bactericcin isolated from a Ca A;Reference number: A58718; MUID:92321788; PMID:1622206
A;Accession: A58718
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                    C;Species: Carnobacterium sp.
C;Date: 23-Jan-1998 #sequence_revision 30-Jan-1998 #text_change 30-Jan-1998
C;Accession: A58718
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h 35.6%; Score 21; DB 2; Le Similarity 44.4%; Pred. No. 6.5e+02; 4; Conservative 3; Mismatches 2;
30.5%; Score 18; DB 2; Liarity 100.0%; Pred. No. 2.8e+05; Conservative 0; Mismatches 0;
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sperm-activating peptide (Tyr-2, Asn-3, Gly-5, Ile-9, Asp-10 SAP-I) - slate-pencil urchi (JSpecies: Heterocentrotus mamillatus (JDate: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 18-Aug-2000 C;Accession: A60589
C;Accession: A60589
C;Accession: A60589
C;Accession: Kajiura, H.; Nomura, K.; Takao, T.; Shimonishi, Y.; Kurita, M.; Yamaguc Comp. Biochem. Physiol. B 94, 739-751, 1989
A;Title: A halogenated amino acid-containing sperm activating peptide and its related peotus mudus, Echinometra mathaei and Heterocentrotus mammillatus.
A;Accession: A60589
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R;Goldschmidt-Reisin, S.; Graack, H.R.
submitted to the Protein Sequence Database, February 1998
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Feb_1998 #sequence_revision 13-Mar-1998 #text_change 21-Jul-2000
                                                                                                                                                                                                                                                                                       C;Date: 18-Jun-1993 #sequence_revision 08-Nov-1996 #text_change 15-Aug-1997 C;Accession: B46453  
R;Takahashi, K.; Kishimoto, S.; Ohori, K.; Yoshizawa, H.; Machida, A.; Ohnuma, H. J. Immunol. 147, 3156-3160, 1991  
A;Title: Molecular heterogeneity of e antigen polypeptides in sera from carriers A;Reference number: A46453; MUID:92013147; PMID:1717588
                                                                                                                                                                                                                                                                                                                                                                                  e antigen p20e precursor - hepatitis B virus N,Alternate names: HBe antigen precursor N;Contains: e antigen C;Species: hepatitis B virus, HBV A,Variety: subtype adr C,Date: 18-Jun-1993 #sequence_revision 08-Nov C;Accession: B46453
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A; Accession: S78420
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| Experimental source: subtype adr
| Experimental source: subtype adr
| Note: sequence extracted from NCBI backbone (NCBIP:60243)
| Note: sequence extracted from NCBI backbone (NCBIP:60243)
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Residues: 1-10 <TAK>
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Pred. No. 2e+03;
1; Mismatches
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Pred. No. 2e+03;
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Pred. No. 2.8e+05;
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C;Accession: I46868
R;Friedman, D.J.; Umeda, P.K.; Sinha, A.M.; Hsu, H. Proc. Natl. Acad. Sci. U.S.A. 81, 3044-3048, 1984
A;Title: Characterization of genomic clones specifying A;Reference number: I46868; NUID:84221901; PMID:6328491
A;Accession: I46868
                                                      R;Ruff-Jamison, S.; Campos-Gonzalez, R.; Glenney Jr., J.F. J. Biol. Chem. 266, 6607-6613, 1991
A;Title: Heavy and light chain variable region sequences A;Reference number: A38740; MUID:91177923; PMID:1706720
A;Accession: A37268
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Accession: S28055
R;Kuntz, M.; Camara, B.; Weil, J.H.; Schantz,
Plant Mol. Biol. 20, 1185-1188, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cytochrome b559 component psbr - pepper chloroplast (fragment)
C;Species: chloroplast Capsicum annuum (pepper)
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_chang
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S28055
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C;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change
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A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-10 <RUF:
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C;Keywords: chloroplast; photosynthesis; photosystem II; thylakoid
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A; Residues: 1-10 < KUN>
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                                                                                                                                                                              Ig heavy chain C region (129) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1997 #sequence_revision 13-Mar-19
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Pred. No. 2.8e+05;
1; Mismatches 1
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C;Species: Mus Sp. (mouse)
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 12-Aug-1996
C;Accession: A47364
R;Shida, M.M.; Ng, Y.K.; Soares, M.J.; Linzer, D.I.
Mol. Endocrinol. 7, 181-188, 1993
A;Title: Trophoblast-specific transcription from the mouse placental lactoge A;Accession: A47364
MUID: 93225959; PMID: 8469232
A;Accession: A47364
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S25266
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R;Serafini, T.; Stenbeck, G.; Brecht, A.; Lottspeich, F.; Orc Nature 349, 215-220, 1991
A;Title: A coat subunit of Golgi-derived non-clathrin-coated A;Reference number: $13636; MUID:91101693; PMID:1898984
A;Accession: $13636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              coat protein beta chain, Golgi-derived - rabbit (fr N;Alternate names: beta-COP protein C;Species: Oryctolagus cuniculus (domestic rabbit) C;Date: 19-Mar-1997 #sequence_revision 24-Mar-1999 C;Date: 19-Mar-1997 #sequence_revision 24-Mar-1999 C;Date: 19-Mar-1999 #sequence_revision 24-Mar-1999 
pilE protein - Escherichia coli (fragment)
C;Species: Escherichia coli
C;Species: Escherichia coli
C;Date: 28-May-1993 #sequence_revision:28-May-1993 #text_change 11-Jun-1993
C;Accession: $25266
R;Dupuy, B.; Taha, M.K.; Possot, O.; Marchal, C.; Pugsley, A.P.
Mol. Microbiol. 6, 1887-1894, 1992
Mol. Microbiol. 6, 1887-1894, 1992
A;Title: PulO, a component of the pullulanase secretion pathway of Klebsiell
A;Reference number: $25266; MUID:92374839; PMID:1354833
A;Accession: $25266
A;Molecule type: protein
A;Residues: 1-7 < DUP>
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Pred. No. 2.8e
3; Mismatches
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Pred. No. 3.1e+03;
1; Mismatches 4;
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2.8e+05;
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PW0002
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A;Molecule type: protein
A;Residues: 1-7 eHI>
C;Keywords: phosphotransferase
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                                                                                 N-methylpurine DNA glycosylase - mouse (fragment) (;Species: Mus musculits (house mouse) (;Date: 19-Jan-1996 #sequence_revision 19-Jan-199 (;Accession: A5602) (R;Roy, R.; Brooks, C.; Mitra, S.
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        R;Roy, R.; Brooks, C.; Mitra, S.
Blochemistry 33, 15131-15140, 1994
A;Title: Purification and biochemical characterization of recombinant N-methylpurine-DNA
A;Reference number: A56029; MUID:95092772; PMID:7999773
                                                                                                                                                                                                                                                      RESULT 14
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polyphosphate-glucose phosphotransferase (EC 2.7.1.63) - Propionibacterium freudenreichi. C.Species: Propionibacterium freudenreichi subsp. shermanii (.7.5) - Propioni S29735 - Propionibacterium freudenreichi subsp. shermanii (.7.5) - Propionibacte
                                                                                                                                                                                                                                                                                                                                                                                                                                           R:LaRoche, J.; Bennett, J.; Falkowski, P.G.
Gene 95, 165-171, 1990
A;Title: Characterization of a cDNA encoding for the 28.5-kDa
A;Reference number: JW0040; MUID:91065528; PMID:2249775
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             chlorophyll a/b-binding protein 31% - green alga (D N;Alternate names: photosystem II light-harvesting C;Species: Dunalisella tertiolecta C;Date: 24-Nov-1999 #sequence_revision 24-Nov-1999
                                                                                                                                                                                                   C;Superfamily: chlorophyll a/b-binding protein
C;Keywords: chloroplast; grana; light-harvesting
                                                    Score 15;
Pred. No.
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Pred. No. 2.8e+05;
l; Mismatches 0;
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ng chlorophyll 31K protein
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#text_change 19-Jan-1996

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hypothetical protein 2 LRH-1 5'-region - mouse
C:Species: Mus musculus (house mouse)
C:Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 18-Aug-2000
C:Accession: S27873
R;Tugwood, J.D.; Issemann, I.; Green, S.
submitted to the EMBL Data Library, February 1992
A:Description: LRH-1: A nuclear hormone receptor active in the absence of exogenous ligal A;Reference number: S27873
A;Recession: S27873
A;Accession: S27873
A;Accession: S27873
A;Accession: S27873
A;Accession: S27873
A;Gross-references: EMBL:M81385; NID:g198872; PIDN:AAA39446.1; PID:g198874
C;Superfamily: unassigned leader peptides
Search completed: May 18, 2004, 10:05:40 Job time: 11.25 secs
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A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-9 <ROY>
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59
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RS19_TOBBP
LCK4_LEUMA
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Matches 4
                                                  Chloroplast.
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
lamiids; Solanales; Solanaceae; Capsicum.
NCBI_TaxID=4072;
 SEQUENCE FROM N.A.
STRAIN=cv. Lamuyo; TISSUE=Fruit, and Leaf;
MEDLINE=93099270; PubMed=1463853;
Kuntz M., Camara B., Weil J.-H., Schantz R.
                                                                                                                                                              (Fragment).
PSBF.
                                                                                                                                                                                        PSBF_CAPAN STANDARD; PRT; 10 AA. (203367; 01-JUN-1994 (Rel. 29, Created) 01-JUN-1994 (Rel. 29, Last sequence update) 10-OCT-2003 (Rel. 42, Last annotation update) Cytochrome b559 beta subunit (PSII reaction c
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P36960;
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NCBI_TaxID=35782;
                                                                                                                                                apsicum annuum (Bell pepper).
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bacte	P81672 pinus pinas	P58807 conus marmo		Q46464 campylobact	P16353 heliothis z	P18110 romalea mic	P80678 chelyosoma	P81749 clupea pall	P02728 homo sapien	P82075 litoria rub	P19343 papio hamad

ALIGNMENTS

01-UUN-1994 (Rel. 29, Created) 01-UUN-1994 (Rel. 29, Last sequence u) 16-OCT-2001 (Rel. 40, Last annotation Lantibiotic carnocin UI49 (Fragment). Carnobacterium sp. (strain UI49). a Carnobacterium sp."; Appl. Environ. Microbiol. 58:1417-1422(1992). -i- FUNCTION: Lanthionine-containing peptide Active on Gram-positive bacteria. Antibiotic; Bacteriocin; Lantibiotic. MEDLINE=92321768; PubMed=1622206; Stoffels G., Nissen-Meyer J., Gud Carnobacterium sp. (strain U149). Bacteria, Firmicutes; Lactobacillales; Carnobacteriaceae; "Purification and characterization of a new bacteriocin isolated from 30.5%; Pro 100.0%; Pro 741776D05B05B810 CRC64; Score 18; DB 1; LC Pred. No. 1.4e+05; Minmarches 0; Gudmundsdottir A., PRT; update) on update) 9 A antibiotic (lantibiotic) Length 7; Sletten Indels ζ. 0 Holo Gaps H

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                                                                                                                                                                                                                                                                                     Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Winstan Davey M., East P.D., Thorpe A.;
"Lepidopteran peptides of the allatostatin superfamily.";
Peptides 18:1301-1309(1997).
-!- SIMILARITY: Belongs to the allatostatin family.
Neuropeptide; Amidation.
Neuropeptide; Amidation.
MOD_RES 8 8 AMIDATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
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Cydia pomonella (Codling moth)
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-!- FUNCTION: This b-type cytochrome is tightly associated with the reaction center of photosystem II and possibly is part of the water-oxidation complex.
-!- SUBUNIT: Heterodimer of an alpha subunit and a beta subunit.
-!- SIMILARITY: Belongs to the psb5 / psbF family.
                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE=Larva;
MEDLINE=98054539; PubMed=9392829;
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InterPro; IPR006216; Cyt_b559.
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                                                                                                                                                                                                                                                            AMIDATION.
0B2879C45B573767 CRC64;
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Pred. No. 1.4e
2; Mismatches
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LUMENAL (POTENTIAL).
817D0F59D6D69DC5 CRC64;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Scott A.G., Winstanley D.,
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  NON TER
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P35946;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NON TER
SEQUENCE
                                                                                EMBL; D10371; BAA01208.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PHODV
                              Transferase; RNA-directed RNA
NON_TER 10 10
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(JUL-1997)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STANDARD;
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RESULT

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1105 MW;

9C2B7FD452D5A2D5 CRC64

polymerase

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USS_FUSSO

D_FUSS_FUSSO STANDARD; PRT; 8 AA.

C_P810IO;
T_15-UTL-1998 (Rel. 36, Created)
T_10-CCT-2003 (Rel. 36, Last sequence update)
T_10-CCT-2003 (Rel. 42, Last annotation update)
T_10-CCT-2003 (Rel. 42, Last annotation update)
E_Allergen Fus s_13596* (Fragment).
S_Pusarium solani (subsp. pisi) (Nectria haematococca).
S_Pusarium solani (subsp. pisi) (Nectria haematococca).
C_Bukaryota; Fungi; Ascomycota; Pezizomycottina; Sordariomy
C_Hypocreomycetidae; Hypocreales; Nectriaceae; Nectria.

K_NCBI_TaxID=70791;
                                                                                                                                                                                                                                                                                                                   STRAIN=Ulster/88;

MEDLINE=9226897; PubMed=1588321;

Curran M.D., O'Loan D., Kennedy S., Rima B.K.;

Curran M.D., O'Loan D., Kennedy S., Rima B.K.;

"Molecular characterization of phocine distemper virus: gene order and sequence of the gene encoding the attachment (H) protein.";

J. Gen. Virol. 73:1199-1194(1992).

-I- FUNCTION: Probable component of the active polymerase. It may function in mRNA synthesis, capping, methylation and poly(A) synthesis of newly synthesized viral mRNAs, RNA editing of the gene transcript, and protein kinase activity.

-I- CATALYTIC ACTIVITY: N nucleoside triphosphate = N diphosphate + (RNA)(N).
This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUN-1994 (Rel. 29, Created)
01-JUN-1994 (Rel. 29, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
RNA polymerase beta subunit (EC 2.7.7.48) (Last protein) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Phocine distemper virus (PDV).
Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.
                                                                                                                                                                                                                                                                                 -!- SIMILARITY: Belongs to the paramyxoviruses L protein family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 L-1997) to Swiss-Prot.
Causes an allergic reaction in human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TISSUE=Mycelium;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    27.1%;
40.0%;
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EFG_CLOPA
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Best Local S
Matches 4
                                                                                                                                                                                        VKI PERAM STANDARD; PRT; 11 AA.

1 PVKI PRT; 11 AA.

1 PVKI PERAM STANDARD; PRT; 11 AA.

1 PVKI PRT; 11 AA.

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Best Local S
Matches 2
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InterPro; IPR000795; EF GTPbind.
PROSITE; PS00301; EFACTOR GTP; PARTIAL.
Blongarion factor; Protein biosynthesis;
NON_TER 11 11
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P81350;
                     SEQUENCE, AND SYNTHESIS.
TISSUE-Abdominal perisympathetic organs;
MEDIINE-95212021; pubMed=7716075;
MEDIINE-95212021; pubMed=7716075;
Perdel R., Linde D., Rapus J., Vettermann S., Penzlin H.;
"Periviscerokinin (Pea-PVK): a novel myotropic neuropeptide
perisympathetic organs of the American cockroach.";
Peptides 16:61-66(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Two-dimensional gel electrophoresis separation and N-terminal sequence analysis of proteins from Clostridium pasteurianum W5. Electrophoresis 19:802-806(1998).

-i- FUNCTION: This protein promotes the GTP-dependent translocathe nascent protein chain from the A-site to the P-site of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=W5;
MEDLINE=98291870; PubMed=9629918;
Flengsrud R., Skjeldal L.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   15-JUL-1998 (Rel
15-JUL-1998 (Rel
28-FEB-2003 (Rel.
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Bacteria; Firmicutes; Cl
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ທ
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       MYOACTIVE
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. 36, Last
. 41, Last
r G (EF-G)
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  PEPTIDE;
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Pred. No. 3e+0
2; Mismatches
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    HAS EXCITORY ACTIONS ON
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3;
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Best Local S
Matches 3
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Best Local S
Matches 3
                     TRPI PSEPU
P36414;
P36414;
O1-JUN-1994
O1-JUN-1994
15-MAR-2004
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P31031;
    01-JUN-1994 (Rel. 29, Created)
01-JUN-1994 (Rel. 29, Last sequence update)
15-MAR-2004 (Rel. 43, Last annotation update)
HTH-type transcriptional regulator trpI (TrpBA operon transcriptional
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; L04163; AAA25660.1; -. PIR; S35490; S35490. REBASE; 1541; PvuI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
STRALN=ATCC 13315;
MEDLINE=93087186;
Smith M.D., Longo M., Gerard G.F.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
10-CCT-2003 (Rel. 42, Last annotation update)
Type II restriction enzyme PvuI (EC 3.1.21.4)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entities requires a license agreement (See or send an email to license@isb-sib.ch).
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Enterobacteriaceae; Proteo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PROVU
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                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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1; Mismatches
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Pred. No. 5
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on it use by non-profit institutions as long as its content is in no was modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Cloning and characterization of genes for the PvuI restriction and modification system.",
Nucleic Acids Res. 20:5743-5747(1992).
-!- FUNCTION: RECOGNIZES THE DOUBLE-STRANDED SEQUENCE CGATCG AND CLEAVES AFTER T-4.
-!- CATALYRIC ACTUVITY: Endonucleolytic cleavage of DNA to give specific double-stranded fragments with terminal 5'-phosphates.
                                                                                                                                          Restriction system; Hydrolase; Muclease; Endonuclease NON_TER 1 1
                                                                                                              9F0CDE7955B72B1A CRC64;
                                                                                                                                                                                                                                                                                                                                         http://www.isb-sib.
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                                                    Length 11;
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P82158;
30-MAY-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
STRAIN-PPG1 C1S;
MEDLINE=89315826; PubMed=2503057;
Eberly L., Crawford I.P.;
"DNA sequence of the tryptophan synthase
                                                                    -!- SIMILARITY: Belongs to the allatostatin
Neuropeptide; Amidation.
MOD_RES 7 7 AMIDATION.
                                                                                                       Duve H., Johnsen A.H., Maestro J.-L., Scott Davey M., East P.D., Thorpe A., "Lepidopteran peptides of the allatostatin Peptides 18:1301-1309(1997).
                                                                                                                                                                                                                                                                          Cydia pomonella (Codling moth).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
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PROSITE; PS50931; HTH_LYSR; PARTIAL
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                                                                                                                                                                                         MEDLINE=98054539; PubMed=9392829;
                                                                                                                                                                                                             TISSUE=Larva;
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Bacteria; Proteoba
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TRPI.
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SIMILARITY: Contains 1 HTH lysr-type DNA-binding domain.
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(Rel. 39, Last sequence update)
(Rel. 39, Last annotation update)
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2; Mismatches
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                                                                                                                                                                          Scott A.G., Winstanley D.,
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                  DB 1;
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Cydiastatin 1.
Cydia pomonella (Codling moth).
Cydia pomonella (Codling moth).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Morricoidea; Tortricidae; Olethreutinae; Cydia.
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P82152;
30-MAY-2000
30-MAY-2000
30-MAY-2000
                                                                                                                                                                                     MOD_RES
SEQUENCE
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Duve H., Johnsen A.H., Macetro J.-L., Scott Davey M., East P.D., Thorpe A.;
"Lepidopteran peptides of the allatostatin s Peptides 18:1301-1309(1997).
-!-SIMILARITY: Belongs to the allatostatin Neuropeptide; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            P81821;
30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Neuropeptide; Amidation; Multigene family MOD_RES 8 AMIDATION (PSQUENCE 8 AA; 919 MW; C82879D5AB569A
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Eumalacostraca; Eucarida; Decapoda; Pleocyemata;
Eubrachyura; Portunoidea; Portunidae; Carcinus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE.
TISSUE=Larva,
                                                                                                                                                                                                                        Neuropeptide;
MOD_RES
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                                                                                                                                                                                        934 MW;
                                                                                 23.7%;
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                                                                                                                                                                                        C82879C45B51F775 CRC64;
                                                 Score 14; DB
Pred. No. 1.4e
1; Mismatches
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Pred. No. 1.4e
2; Mismatches
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1.4e+05;
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UF02_MOUSE
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P83281;
C P83281;
T 28-FEB-2003 (Rel. 41, Created)
T 28-FEB-2003 (Rel. 41, Last sequence update)
T 28-FEB-2003 (Rel. 41, Last annotation update)
E FMRPamide-like neuropeptide FLP8 (VSHNNFLRF-amide).
S Macrobrachium rosenbergii (Giant fresh water prawn).
C Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
C Eumalacostraca; Eucarida; Decapoda; Pleccyemata; Caridea;
C Palaemonoidea; Palaemonidae; Macrobrachium.
N NCBI TaxID-79674;
                Query Match
Best Local (
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Best Local (
   Matches
                                                            Merrick B.A., Patterson R.M., Wichter L.L., He C., Selkirk J.K.;
"Separation and sequencing of familiar and novel murine proteins
using preparative two-dimensional gel electrophoresis.";
Electrophoresis 15:735-745(1994).
-i- MISCELLANEOUS: On the 2D-gel the determined pI of this unknow
protein is: 6.0, its MW is: 32 kDa.

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SEQUENCE 9 AA; 1102 MW; 7E73EAB6D05B1AAB CRC64;
                                                                                                                                                                                                                                                                                   UF02_MOUSE STANDARD; PRT; 9 AA.
P38640;
01-OCT-1994 (Rel. 30, Created)
01-OCT-1994 (Rel. 30, Last sequence update)
15-MAR-2004 (Rel. 43, Last annotation update)
Unknown protein from 2D-page of fibroblasts (P32) (Fragment).
Mus musculus (Mouse).
                                                                                                                                                                                 SEQUENCE.
SEQUENCE.
TISSUB-Fibroblast;
MEDLINE-95009907; PubMed-7523108;
                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Neuropeptide; Amidation.
MOD_RES 9 9
SEQUENCE 9 AA; 1133 M
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-i- MASS SPECTROMETRY: NWW-1133.8; METHOD-MALDI.

-i- SIMILARITY: Belongs to the FARP (FMRFamide related peptide)
                                                                                                                                                                                                                                                NCBI_TaxID=10090;
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Sithigorngul P., Saraithongkum W., Longyant S., Panchan N.,
Sithigorngul W., Petsom A.;
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GO:0007218; P:neuropeptide signaling
h 23.7%; So Similarity 100.0%; I 3; Conservative 0;
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Pred. No. 1.4e+05;
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Matches 2
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1: FUNCTION: May act as a neurotransmitter or neuromodulator.

1: SIMILARITY: Belongs to the allatostatin family.

Neuropeptide; Amidation; Multigene family.

Neuropeptide; Amidation, Multigene family.

Neuropeptide; Amidation, Multigene family.
                                                                                                                                                                                                                                                                                              SEQUENCE
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Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyura;
Eubrachyura; Portunoidea; Portunidae; Carcinus.
NCBI_TaxID=6759;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE=Cerebral ganglion, and Thoracic ganglion MEDLINE=98121193; PubMed=9461295;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Thorpe A.;
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30-MAY-2000 (Rel.
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Pred. No. 7.2e+03;
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3569AB1 CRC64;
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Q1-MAY-2000 (TrEMBLrel. 13, C)
Q1-MAY-2000 (TrEMBLrel. 13, L)
Q1-JUN-2002 (TrEMBLrel. 21, L)
Q1-JUN-2002 (TrEMBLrel. 21, L)
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RESULT
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ID P8
AC P8
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P82081;
01-MAY-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Fibrinolytic metalloproteinase (Fragment).
Agkistrodon contortrix.
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scieroglossa; Serpentes; Colubroidea;
Viperidae; Crotalinae; Agkistrodon.
MCBI_TaxID=8720;
                                                                                                                                                                                                                                                                                                                                                  Guan A.L., Retzios A.D., Henderson G.N., Markland F.S.Jr.; "Purlication and characterization of a fibrinolytic enzyme from venom of the southern copperhead snake (Agkistrodon contortrix contortrix).";
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MEDLINE=91378546; PubMed=1898066;
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ER 11 11
NCE 11 AA; 1209 MW; 7CA02D1D41E87
(TrEMBLrel. 13, (TrEMBLrel. 13,
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71.4%;
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Last annotation updat
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Pred. No. 1.4e+02;
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Last sequence update)
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01-MAY-2000
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DYNASTIN 4
Limnodynastes salmini (Salmin's-striped frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Myobatrachi
Limnodynastinae; Limnodynastes.
NCBI TaxID=39404;
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Aust. J. Chem. 46:833-842(1993).
-!- MASS SPECTROMETRY: MW=1236; METHOD=FAB.
SEQUENCE 12 AA; 1236 MW; 147AA70FDF472724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bradford A.M., Raftery M.J., Bowie J.H., Wallace J.C., "Peptides from Australian frogs. The structure of the Limnodynastes salmini and Fletcherin from Limnodynaste Aust. J. Chem. 46:1235-1244(1993).

-!- MASS SPECTROMETRY: MW=772; METHOD=FAB.
SEQUENCE 8 AA; 772 MW; 7B58772455A2C728 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Limnodynastes salmini (Salmin's-striped frog).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Myobatrachidae;
Limnodynastinae; Limnodynastes.
                                                                                                                                                                                                            DYNASTIN
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                                                                                                                                                                                                                                                                                                                                                                                            P82083
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Eukaryota; Metazoa; Chordata; Craniata; Ve
Amphibia, Batrachia; Anura; Neobatrachia;
Limnodynastinae; Limnodynastes.
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DYNASTIN 3.
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Pred. No. 1e+0
1; Mismatches
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Pred. No. le+03;
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ta; Vertebrata; Euteleostomi;
chia; Bufonoidea; Myobatrachidae;
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                                                                                                            Bufonoidea; Myobatrachidae;
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  odynastes fletcheri.";
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P82085;
01-MAY-2000 (TrEMBLrel 13, C
01-MAY-2000 (TrEMBLrel 13, I
01-MAY-2000 (TrEMBLrel 13, I
DYNASTIN 7.
                                                                                                                                                                                                                                                                                                          Q34909;
Q34909;
01-NOV-1996
01-NOV-1996
01-JUN-2003
SEQUENCE FROM N.A.

MEDLINE=8823478; PubMed=2836084;

MCCracken A., Uhlenbusch I., Gellissen G.;

"Structure of the cloned Locusta migratori:

restriction mapping and sequence of its ND

Curr. Genet. 11:625-630(1987).

EMBL; X05286; CAA28905.1; -.

GO; GO:0005739; C:mitochondrion; IEA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE, AND MASS SPECTROMETRY.
TISSUB-SKIN SECRETION;
TISSUB-SKIN SECRETION;
Bradford A.M., Raftery M.J., Bowie J.H., Wallace J.C., Tyler M.J.;
"Peptides from Australian frogs. The structure of the dynastins from
"Peptides from Australian frogs. The structure of the dynastins from
Limnodynastes salmini and Fletcherin from Limnodynastes fletcheri.";
Aust. J. Chem. 46:1235-1244(1993).
-!- WASS SPECTROMETRY: MW=1114; METHOD=FAB.
SEQUENCE 12 AA; 1114 MW; 3AB5A976CAA72728 CRC64;
                                                                                                                                                                                           Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridomorpha;
Acridoidea; Acrididae; Oedipodinae; Locusta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Limnodynastes salmini (Salmin's-striped fi
Bukaryota; Metazoa; Chordata; Cranista; Ve
Amphibia; Batrachia; Anura; Neobatrachia;
Limnodynastinae; Limnodynastes.
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-!- MASS SPECTROMETRY: MW=786; METHOD=FAB.
SEQUENCE B AA; 786 MW; 7B58772455B05728 CRC64;
                                                                                                                                                                    NCBI_TaxID=7004;
                                                                                                                                                                                                                                                        Mitochondrion.
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Pred. No. 4.3e
1; Mismatches
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Vertebrata; Euteleostomi;
a; Bufonoidea; Myobatrachidae;
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4.3e+03;
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Best Local Similarity
Matches 4; Conserv
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071066;
071066;
01-AUG-1998 (TrEMBLrel. 07, C
01-AUG-1998 (TrEMBLrel. 07, L
01-DEC-2001 (TrEMBLrel. 19, L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        092766
092766;
01-NOV-1998
01-NOV-1998
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NON_TER 1

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SEQUENCE 8 AF
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NON TER
SEQUENCE
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Liermann H., Harder T., Haas L.;
"Genetic analysis of the central un
proximal coding part of the F gene
morbilliviruses.";
Submitted (SEP-1997) to the EMBL/Ge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Viruses; ssRNĀ negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.
NCBI_TaxID=11232;
                                                                                                                                                                                                                                                                           Canine distemper virus.
Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.
                                                   Liermann H., Harder T., Haas L.;
"Genetic analysis of the central untranslated genome region proximal coding part of the F gene of wild-type and vaccine morbilliviruses.";
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
EMBL; AF028234; AAC00164.1; -.
                                                                                                                                                                      SEQUENCE FROM N.A.
STRAIN=Dog #10757/96;
Liermann H., Harder T
                                                                                                                                                                                                                                                      Paramyxoviridae;
NCBI_TaxID=11232;
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                     SEQUENCE
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                                          AF026234;
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ER 9 9
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3; Conserv
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B (TrEMBLrel. 08,
8 (TrEMBLrel. 08,
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ilarity 100.0%;
Conservative (
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                     1124 MW;
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0; Mismatches
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Pred. No. le+06;
2; Mismatches
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Matches 3
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Best Local S
Matches 2
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Best Local Similarity
Matches 3; Conser
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SEQÜENCE 10 AA.
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Q9PXC3;
Q9PXC3;
Q1-MAY-2000 (TrEMBLrel. 13, C
01-MAY-2000 (TrEMBLrel. 13, I
01-UN-2003 (TrEMBLrel. 24, I
                                                                                                                                                                                                                MEDLINE=92013147; PubMed=1717588;
Takahashi K., Kishimoto S., Ohori K., Yoshizawa H.,
Takahashi K., Tsuda F., Munekata E., Miyakawa Y., Mayum
Ohnuma H., Tsuda F., Munekata E., Miyakawa Y., Mayum
"Molecular heterogeneity of e antigen polypeptides i
carriers of hepatitis B virus.";
J. Immunol. 147:3156-3160(1991).
                                                                                                                                                                                                                                                                                                                                                                                         B antigen P20E (Fragment).

Heparitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407;
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Bukaryota; Viridiplantae; Stra
Spermatophyta; Magnollophyta;
Ehrhartoideae; Oryzeae; Oryza
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01-NOV-1996 (TrEMBLrel 01,
01-NOV-1996 (TrEMBLrel 01,
01-OCT-2002 (TrEMBLrel 22,
Alpha-amylase (Fragment).
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NON_TER 8
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EMBL; M62916; AAA33892.1;
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Rodriquez R.L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=91078641; PubMed=2258052;
Kumagai M.H., Shah M., Terashima M.,
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6 HWTOI S
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Conservative 0;
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                                                                                                                                                        1238 MW;
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yta; Liliopsida;
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0; Mismatches
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1; Mismatches
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Pred. No. 1e+0
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                                                                                                                                                                                                                                                                  akawa Y., Mayumi M.;
polypeptides in sera from
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                                                                 DB 12; I
9.1e+03;
thes 1;
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1e+06;
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Indels

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Gaps

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Indels

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Machida A.,

Whitaker J.R.,

saccharomycers

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RESULT
Q23876
ID
Q23876
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Q23876
Q22876
Q22976
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Q28742
ID Q2874
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Q85631;
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01-NOV-1996
01-DEC-2001
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Q23876;
Q1-NOV-1996
Q1-NOV-1996
Q1-NOV-1998
Q28742 PRELIMINARY; PRT; 7 AA. Q28742; Q28742; 01-NOV-1996 (TrEMBLrel. 01, Created) 01-NOV-1996 (TrEMBLrel. 01, Last sequence update) 01-JUN-2003 (TrEMBLrel. 24, Last annotation update) Alpha-myosin heavy chain (Fragment). Oryctolagus cuniculus (Rabbit).
                                                                                                                                                                                                                                                                                                                                    EMBL; K03
NON TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=82260445; PubMed=6286214;
McKeown M., Firtel R.A.;
"Actin multigene family of Dictyostelium.";
Cold Spring Harb. Symp. Quant. Biol. 46:495-505(1982)
EMBL; K02957; AAA33150.1; -.
EMBL; K02956; AAA33150.1; JOINED.
                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=85033920; PubMed=6092695;
Sutrave P., Jansen H.W., Bister K., Rapp U.R.;
"3'-terminal region of avian carcinoma virus MH2 shares
elements with avian sarcoma viruses Y73 and SR-A.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=11958;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Avian carcinoma virus.
Viruses; Retroid viruses; Retroviridae; Alpharetrovirus.
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Eukaryota; Mycetozoa; Dictyosteliida;
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K03100; AAA42388.1; -.
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G (TrEMBLrel.
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al DNA, myc to 3' LTR (Fragment).
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le+04;
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hes 0;
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Matches 4
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Q9GD12;
Q9GD12;
Q9GD12;
Q9GD12;
O1-MAR-2001 (TEMBLrel. 16, Cr
O1-MAR-2001 (TEMBLrel. 24, L;
O1-UN-2003 (TEMBLrel. 24, L;
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Best I
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Q15891;
01-NOV-1996 (TREMBLrel. 01, 0
01-NOV-1996 (TREMBLrel. 01, 1
01-DEC-2001 (TREMBLrel. 19, 1
(Clone XP2E8B) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                   "Isolation of chromosome-specific genes arrayed cDNAs and cosmid libraries."; Hum. Mol. Genet. 0:0-0(1995).
EMBL; L32131; AAA73881.1; -.
NON_TER 1 1
NON_TER 9 9
SEQUENCE 9 AA; 1030 MW; E56635A1A336
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens (Human)
Eukaryota; Metazoa;
Mammalia; Eutheria;
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SEQUENCE FROM N.A.
MEDLINE=84221901; PubMed=6328491;
MEDLINE=84221901; PubMed=6328491;
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NON_TER
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          Chloroplast.
Eukaryota; Viridiplantae; Streptophyta; Er
Spermatophyta; Magnoliophyta; Liliopsida;
                                                                               RPS16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TISSUE=Placenta;
Lee C.-C., Yazdani A., We
Lee C.-C., Yazdani A., We
Coolbaugh M.I., Chinault
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Mammalia; Eutheria;
                                                                 Linospadix monostachya.
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3ci. U.S.A. 81:3044-3048(1984)
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Primates;
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lt C.A.,
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Baldini A., Lindsay E.A.,
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              Embryophyta;
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                Tracheophyta, Arecoideae;
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OC Areceae; Linospadicinae; Linospadix.

OX NCBI TaxID=131282;

RN [1]

RN [2]

RN [1]

RP SEQUENCE FROM N.A.

RC TISSUE-Leaf;

RA Asmussen C.B., Chase M.W.;

RI "Coding and noncoding plastid DNA in palm systematics.";

RI Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.

RL Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.

DR GO: GO:0009507; C:chloroplast; IEA.

Chloroplast.

I 1

FT NON_TER 1

FT NON_TER 9

SQ SEQUENCE 9 AA; 1106 MW; 1B9236C2C0441B50 CRC64;

Query Match 28.8%; Score 17; DB 8; Length 9;

Best Local Similarity 75.0%; Pred. No. 1e+06;
Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Cy 5 IQLM 8

Db 6 VQLM 9

Search completed: May 18, 2004, 10:04:47

Job time : 32.25 secs
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Maximum Match 100%
Listing first 45 summaries
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Copyright (c) 1993 - 2004 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		•			SUMMARIES	
Result	Score	Query Match	Length	BG	ID	Description
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7	49	83.1	10	σ	16	66 Human
89	45	٥,	9	4	AAB86225	25 Human
9	43	Ν.	9	N	AAR91645	5 Human
10	43	'n	9	w	AAY96981	31 Parati
11	43	'n	9	4	AAB86220	õ
12	40.5	68.6	11	N	AAY50600	0 Resin
13	40	.7	9	w	AAB01862	
14	40	.7	9	w	AAY96966	Aay96966 Parathyro
15		<i>∾</i>	9	ω	AAB01863	•
16		2	10	4.	AAB96932	Aab96932 Rat parat
17		2	11	4.	AAB96931	Aab96931 Rat parat
18		2	11	4	AAB96915	Aab96915 Parathyro
19		62.7	11	4	AAB84770	Aab84770 Parathyro
20		.2	12	4.	AAB96914	Aab96914 Parathyro
21		۶	12	4.	AAB84769	
22		61.0	9	ω	AAY78849	ø
23	<u>ა</u>	٠	. 00	N	AAR91646	9
24		59.3	00	ω	AAB07467	Aab07467 Antigenic
25		59.3	8	4	AAB86221	2

CCCCCCCCCXXXXIIIXX

Claim 4; Page 92; 119pp; English.

Compounds of the structure or formula S-(L)_n-B, R1-S-(L)_n-R or S-(L)_n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-rP) (1-34); R_1 is the PTH-1 receptor signal sequence; and R is (a portion of) PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAMP

New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.

WPI; 2000-452384/39.

<u>4</u> 5	44	43	42	41	40	39	38	37	36	ა წ	3 4	33	32	3 <u>1</u>	30	29	28	27	26
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76	AAR91648	AAB21270	AAR89301	ABU54029	AAU93357	ABG69386	9	AAB01864	N	AAR91647	Q	AAB96892	AAP82547	σı,	ABP71482	AAW45785	ABP71483	ABP71485	ABP71484
Aay68764	Aar91648	Aab21270	Aar89301	Abu54029	Aau93357	Ω.	Aay97062	Aab01864	Aab86222	Aar91647	Aab96891		Aap82547	Aab01866	ထ	Aaw45785	\vdash	Abp71485	ADD/1484
Amino aci	Human par	Fertilin	Japanese	Human DNA	Granulocy	Vascular	PTH-rP N-	PTH(1-14)	Human par	Human par	Rat parat	Rat parat	(Asn10, T	PTH(1-14)	Parathyro	Parathyro	Parathyro	Parathyro	Paracnyro

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Gardella TJ, Kronenberg HM, Potts JT, Jueppner H;		_	(POTT/) POTTS J T.	_	(GARD/) GARDELLA T J.		31-DEC-1998; 98US-0114577P.	30-DEC-1999; 99WO-US031108.		(06-JUL-2000.)		WC200039278-A2.		Homo gapiens		reformation; resorption; remodeling; tether1; osteoporo:	PTH; parathyroid hormone; N-terminal; signaling domain; bone mass;		Parathyroid hormone N-terminal signaling domain (residues 1-11).		31-OCT-2000 (first entry)		AAY96968;		AAY96968 standard; peptide; 11 AA.	RESULT 1

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RESULT 2
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Best Local &
The invention relates to a new antigenic peptide for inducing the formation and isolation of antibodies having an affinity to it, being formed from the N-terminus of parathyroid hormone (PTH). Also included are; (1) a method for producing antibodies useful in the determination PTH levels in a biological sample comprising; (a) providing at least on first peptide antigen comprising a peptide fragment of PTH; (b) administering the first peptide antigen to a host animal to induce antibody production; (c) monitoring the antibody titre produced; (d) isolating antisera produced in the host animal; and (e) selecting antisera from the isolated antisera produced in the host that is capable of binding to a second peptide antisera; (2) an antibody (ab) produced be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bos taurus.
Sus scrofa.
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                                                                                                                                                                                                                                                                                                                                                                                                        New parathyroid hormone (PTH) antigenic peptide inducing the and isolation of antibodies having an affinity to it, useful determining bioactive PTH levels in serum, plasma and/or cell
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LAVIGNE J R.
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RESULT 3
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                                                                                                       Zahradnik RJ,
                                                                                                                             (ZAHR/)
(LAVI/)
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LAVIGNE J R.
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The invention relates to a new antigenic peptide for inducing the CC formation and isolation of antibodies having an affinity to it, being CC are; (1) a method for producing antibodies useful in the determination of CPTH levels in a biological sample comprising: (a) providing at least one first peptide antigen comprising a peptide fragment of PTH; (b) antibody production; (c) monitoring the antibody titre produced; (d) CC antibody production; (c) monitoring the antibody titre produced; (d) CC isolating antisera produced in the host animal to induce CC antisera from the isolated antisera produced in the host that is capable CC isolating and (3) test kits and analytical procedures used for the CC determination of bioactive intact PTH utilising (ab). The methods and CC compositions of the present invention are useful for determining CC disolative intact PTH levels in serum, plasma and/or cell culture media. CC compared to prior art, have the particular advantages of possessing CC compared to prior art, have the particular advantages of possessing CC compared to prior art, have the particular advantages of possessing CC prients suffering from hypercalcaemia, osceporosis and primary CC hyperparathyridism. The present sequence represent a PTH antigenic peptide representing amino acids 1-12 of human, mouse, rat, porcine, CC canine and bovine PTH

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RESULT 4
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                                                                                                                New antigenic peptide(s) from human parathyroid hormone - and antibodies generated using them, able to distinguish between active and inactive forms of the hormone.
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                                                                                                                                                                                                                                                                                                                                                                                                                                       (FORS/) FORSSMANN W.
                                                                                                                                                                                                                                                                             1996-180391/19
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                                         Page 4; 5pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           active hPTH 1-37.
                                                                                                                                                                                                                                                                                                                                                    Forssmann
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hormone; hPTH; antigen; alpha-helix; immunogen;
                                             German
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                                                                                                                                                                                                                                                                                                                                                        Maegerlein M;
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B S

Matches

10; Similarity

Conservative

0

Mismatches

Indels

0,

Gaps

0

83.1%; 100.0%;

Score 49; ; Pred. No.

DB 3; . 0.017;

Length 10;

1 SVSEIQLMHN 10

Local

Sequence 10 AA;

firming

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RESULT 5
AAY68767
ID AAY6
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Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is a specific example of claimed immunogenic peptides having a sequence from hPTH(1-37) which includes the N-c terminal alpha-helical region and/or the non-structured region of hormone. Antibodies and their binding fragments generated by injection animal with the peptides are useful as diagnostic reagents for determination of biologically active hPTH(1-37)
                                     invention have lipolysis stimulating activity (especially when topically administered). The lipolytic activity of the peptides is enhanced when they are chemically modified to increase their lipophilicity. The peptides are used in cosmetic or dermatological compositions for skin care. They are especially used for slimming treatment of excessive weight in the thighs and hips, in the treatment of cellulite and for skin
                                                                                                                                                                                                   New parathyroid hormone fragment peptides, used as lipolysis stimulants in topically applied cosmetic compositions for slimming treatment of excessive weight in hips and thighs.
                                                                                                                                                                                                                                                                                                                                                                                                            27-JAN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAY68767;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                07-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Parathyroid hormone; lipolysis; cosmetic; de slimming treatment; cellulite; skin firming.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Amino acids 1-10 of a parathyroid hormone (pTH).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAY68767 standard; peptide; 10 AA
                                                                                                                             The present sequence represents a parathyroid hormone (pTH) fragment, comprising amino acids 1-10. Parathyroid hormone fragments of the
                                                                                                                                                                      Claim 1; Page 8; 18pp; French.
                                                                                                                                                                                                                                                             WPI; 2000-171243/15.
                                                                                                                                                                                                                                                                                                                                                    17-JUL-1998;
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                                                                                                                                                                                                                                                                                          Lintner K;
                                                                                                                                                                                                                                                                                                                        (SEDE-) SEDERMA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 Similarity 100
10; Conservative
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Pred. No.
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0.017;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               dermatological; skin care;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 10;
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VSEIQLMHNLG 12

Matches Query Match Best Local

10;

Conservative

Similarity

84.7%; 90.9%;

Score 50; DB 6; Pred. No. 0.014;

Length 12; Indels

Mismatches

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0

0

Sequence 12 AA;

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RESULT 6
RABBG219
ID AABB
XX AABB
XX AABB
XX O3-S
XX Para
XX AABB
XX O3-S
XX D1-D
XX O1-D
XX O
RESULT 7
ABR44166
ID ABR4
XX
AC ABR4
XX
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                                                                                                                                                                                                                                                                                                               S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes a novel method for determining (M1) the content of active parathyroid hormone (A) by treating a sample with (i) antibody (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), and including the N-terminal residue and (ii) antibody (Ab2) that recognizes an epitope within the receptor-binding site of (A). The number of molecules that react with both antibodies is determined and used to calculate the content of physiologically active (A). The method is used for diagnosis of calciun-metabolism disorders, osteopathy, or hypo- or hyper-parathyroidism. The method (unlike known assays) recognizes that (i) some fragments of (A) shorter than the complete (84 aa) peptide are active and (ii) that apparently intact peptide may be biologically cinactive, and also takes into account the fact that some fragments of (A) are antagonistic (these have the receptor-binding site but lack the N-terminus). It thus provides a true measure of the content of physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This sequence represents a peptide fragment used to illustrate the method of
                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibodies reactive with different epitopes.
                              ABR44166
                                                                                          ABR44166 standard; peptide; 10 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 3; 10pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-376318/40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            03-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hypo-parathyroidism; hyper-parathyroidism.
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nes 10; Conserv
                                                                                                                                                                                                                                                                                                               SVSEIQLMHN 10
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100.0%; Pr
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0.017;
hes 0;
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RASULT 8
AAB862
ID AAB8
XX AAB8
AC AAB8
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XX AAB8
XX AAB8
XX Huma
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XX Para
AKW diag
XX Homo
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XX IIIII
XX IIIIII
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Chang M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a fusion peptide (Tat-hPTHDP), where self cell-penetrating Tat peptide is bound to human parathyroid hormone-derived peptide (hPTHDP). The fusion peptide is useful as a component of skin slimming cosmetic composition. The fusion peptide does not cause irritation, easily and safely penetrates into integument and endothelium, does not cause skin disease and has superior lipolysis effects, and is durable. The present sequence represents a human parathyroid hormone (hPTH) fragment that can be used to construct the fusion peptide
                                                                                                                                                                                 Homo
                                                                                                                                                                                                                            Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist; hypo-parathyroidism; hyper-parathyroidism.
                                                                                                                                                                                                                                                                                                                                    Human parathyroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel fusion peptide comprising self to human parathyroid hormone-derived slimming cosmetic composition.
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15-MAR-2002; 2002KR-00014062.
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10; Conserv
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0;
                                                                                                                                                                                                                                                                                                                                    immunogenic peptide SEQ ID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cell-penetrating Tat peptide bound peptide, useful as component of skin
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RESULT 9
AAR91645
ID AAR9
XX AAR9
AC AAR9
XX Huma
XX Huma
XX Huma
XX Huma
XX Huma
XX Huma
XX Of AAR9
XX Synt
XX DE44
XX DE44
XX DE44
XX DE44
XX ACA
PD 04-A
XX Z8-S
XX Z8-S
XX FR Ade1
XX WPI
XX WPI
XX WPI
XX WPI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC This invention describes a novel method for determining (M1) the content CC of active parathyroid hormone (A) by treating a sample with (i) antibody CC (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), CC and including the N-terminal residue and (ii) antibody (Ab2) that CC recognizes an epitope within the receptor-binding site of (A). The number CC of molecules that react with both antibodies is determined and used to CC calculate the content of physiologically active (A). The method is used for diagnosis of calcum-metabolism disorders, osteopathy, or hypo- or CC hyper-parathyroidism. The method (unlike known assays) recognizes that CC (i) some fragments of (A) shorter than the complete (84 aa) peptide are active and (ii) that apparently intact peptide may be biologically CC active and also takes into account the fact that some fragments of (A) care antagonistic (these have the receptor-binding site but lack the N-CC terminus). It thus provides a true measure of the content of physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This sequence represents a peptide fragment used to illustrate the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibodies
                                                                                                                   28-SEP-1994;
                                                                                                                                                     28-SEP-1994;
                                                                                                                                                                                       04-APR-1996
                                                                                                                                                                                                                           DE4434551-A1.
                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                              Human parathyroid
                                                                                                                                                                                                                                                                                                                                                                                        06-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                                                                          AAR91645
                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR91645 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 3; 10pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         reactive with different epitopes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2001-376318/40
                                                                               (FORS/) FORSSMANN W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 invention
                                                                                                                                                                                                                                                                                                                                                  parathyroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       IMMUNDIAGNOSTIK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  VSEIQLMHN 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   VSEIQLMHN 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      76.3%; Score 45; DB 4; Lilarity 100.0%; Pred. No. 1.4e+06; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                 94DE-04434551
                                                                                                                                                     94DE-04434551
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99DE-01061350.
                                                                                                                                                                                                                                                                                                                hormone; hPTH; antigen; alpha-helix; immunogen
                                                                                                                                                                                                                                                                                                                                                  hormone antigenic peptide hPTH 1-9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 9;
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WPI; 1996-180391/19

Adermann K,

Forssmann

Ξ,

Hock D,

Maegerlein M;

Compounds of the structure or formula S-(L) n-B, R 1-S-(L) n-R or S-(L) n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-rP) (1-34); R 1 is the PTH-1 receptor signal sequence; and R is (a portion of) PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAMP in a mammalian cell having PTH-1 receptors, or screening for a peptide or

New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.

WPI; 2000-452384/39

Claim 11; Page 93; 119pp; English.

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AAY96981
ID AAY96
XX AAY96
XX AAY96
XX DT 31-OC
XX PTH;
XW DONE
XX POTH;
XW DONE
XX HOMO
XX WOZOO
XX CGARD
PA (GOTTI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (GARD/) GARDELLA T J
(KRON/) KRONENBERG F
(POTT/) POTTS J T.
(JUEP/) JUEPPNER H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is a specific example of claimed immunogenic peptides having a sequence from hpTH(1-37) which includes the N- or C-terminal alpha-helical region and/or the non-structured region of the hormone. Antibodies and their binding fragments generated by injecting animal with the peptides are useful as diagnostic reagents for determination of biologically active hPTH(1-37)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New antigenic peptide(s) from human parathyroid hormone - and antibodies generated using them, able to distinguish between active and inactive forms of the hormone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gardella TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                31-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                30-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Parathyroid hormone N-terminal signaling domain.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY96981 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; Page 4; 5pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            parathyroid hormone; N-terminal; signaling domain; bone mass; reformation; resorption; remodeling; tether1; osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 SVSEIQLMH 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SVSEIQLMH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kronenberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                98US-0114577P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99WO-US031108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              H 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          72.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , MH
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Pred. No. 1.4e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Potts JT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
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RESULT 11
AAB86220
ID AAB86
XX AAB86
XX AAB86
XX AAB86
XX Parat
DT 03-SE
XX Parat
KW Darat
CS Homo
XX DE199
XX 17-DE
XX 17-DE
XX 17-DE
XX ATMD
XX ATMD
XX Deter
PT usefv
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                                                        This invention describes a novel method for determining (M1) the content of active parathyroid hormone (A) by treating a sample with (i) antibody (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), and including the N-terminal residue and (ii) antibody (Ab2) that recognizes an epitope within the receptor-binding site of (A). The number of molecules that react with both antibodies is determined and used to calculate the content of physiologically active (A). The method is used for diagnosis of calcium-metabolism disorders, osteopathy, or hypo- or hyper-parathyroidism. The method (unlike known assays) recognizes that (i) some fragments of (A) shorter than the complete (84 aa) peptide are active and (ii) that apparently intact peptide may be biologically inactive, and also takes into account the fact that some fragments of (A) contrast methods that some fragments of (A) are antagonistic (these have the receptor-binding site but lack the N-terminus). It thus provides a true measure of the content of and its 1-37 fragment which may produce falsely high values. This sequence represents a peptide fragment used to illustrate the method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibodies reactive with different epitopes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 3; 10pp; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     non-peptide PTH (clinhalation unlike tregular injections
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
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9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                hormone
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Pred. No. 1.4e+06;
0; Mismatches 0;
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This invention describes a novel method for the preparation of peptides (II) that contain both cyclic and linear peptide fragments comprises sequential reaction of a resin-bound linear fragment with the cyclic fragment in N-protected form and optionally other linear fragments. The products of the invention have osteopathic and hypotensive activity. (II) bind to hpTH receptors and act as agonists or antagonists of hPTH. The method is particularly used to prepare cyclic peptide analogs of parathyroid hormone (PTH) or PTH-related peptides which are useful for treating diseases that respond to treatment with agents that bind to PTH receptors (with or without activation of adenylyl cyclase activity), e.g. hyper- or hypo-calcemia, osteoporosis, osteopenia, hyper- or hypo-calcemia, cushing's syndrome, renal failure and hypertension, also for promoting repair of bone fractures. Separate synthesis of the cyclic fragment allows convergent synthesis of resin-bound (II), with better yields and higher throughput. The difficulties associated with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 75; 85pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Convergent synthesis of peptides for treating e.g. bone disorders
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1999-633822/54
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The invention relates to a novel parathyroid hormone (PTH) peptide (AAB01853) and parathyroid hormone-related peptide (PTHP); AAB01860), and biologically active derivatives thereof (AAB01857-B01858, AAB01861).

B01869). The peptides of the invention are at least 85% identical to the generic peptide of the formula: XI-Val-Ser-Glu-X2-Gln-Leu-X3-His-X4-X5-CC Gly-Lys-X6 (AAB01857) where: X1 is Ser or Ala; X2 is 11e or Lys; X3 is CC Met, Leu or Nle; X4 is Asn or Asp; X5 is Leu or Lys; X6 is His or Ser; provided that the peptide is not PTHP(1-14). The peptides of the invention consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C-cterminal derivatives. PTH is a major regulator of calcium homeostasis, and is necessary for the normal function of the gastrointestinal, skeletal, neurological system, neuromuscular and cardiovascular systems. It binds to both PTH-1 receptors on osteoblasts and renal tubular cells, and to the recently identified PTH-2 receptor. PTH has a potent anabolic effect on the skeleton, and mediates calcium reabsorption, enhances of the renal and skeletal actions of PTH, and also bind to the PTH-1
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Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Parathyroid hormone peptide; PTH; PTH-related peptide; PTHrP; calcium homeostasis; PTH-1 receptor; PTH-2; vitamin D synthesis; bone synthesis; agonist; osteoporosis; non-parenteral delivery.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                                                                                                                 Disclosure; Page 26; 73pp; English.
                                                                                                                                                                                                                                                                                                                                             Parathyroid hormone (PTH) peptides, PTH related peptides and the nucleic acids that encode them, useful for treating osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GARD/)
(KRON/)
(POTT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          can be purified before reaction with the resin-bound component. AAY50568-Y50614 represent the peptide fragments described in the method of the
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                                                          receptor. They do not bind to the PTH-2 receptor. The peptides of the invention are either agonists of pTH-1 and pTH-2 receptors (AAB01858, AAB01861-B01866) or are PTH-1/PTH-2 receptor agonists (AAB01867-B01869). PTH-1/PTH-2 receptor agonists are useful for the treatment of conditions characterised by a decrease in bone mass, such as osteoporosis. PTH-1/PTH-2 receptor antagonists are useful for treatment of some particular action of the PTH-1/PTH-2 receptor antagonists are useful for treating medical disorders that arise from excessive or altered action of the PTH-1/PTH-2 receptor. Detectably labelled peptides of the invention are also useful in the determination of rates of bone formation, bone resorption and/or bone remodelling in a patient. The peptides of the invention are "minimised" versions of PTH or PTH-P which are inexpensive to prepare by conventional synthetic chemistry, and can be delivered to a patient via non-parenteral routes. Sequences AAB01858 and AAB01861-B01866 represent PTH-1/PTH-2 receptor agonists
Sequence 9
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PTH;
WO200039278-A2
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                                                                                                                                                                                                                                   AAY96966;
                                                                                                                                                                                                                                                                          AAY96966 standard; peptide; 9
                                          Homo sapiens
                                                                                                                                              Parathyroid hormone N-terminal signaling domain (residues 1-9).
                                                                              parathyroid hormone; N-terminal; signaling reformation; resorption; remodeling; tether
                                                                                                                                                                                         (first entry)
                                                                                     tether1;
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31-DEC-1998;
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                                               06-JUL-2000
 98US-0114577P
                       99WO-US031108
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Gardella TJ, Kronenberg , MH Potts JT, Jueppner

(KRON/)

) KRONENBERG H M.) POTTS J T.) JUEPPNER H.

(GARD/)

GARDELLA

(JUEP/)

2000-452384/39

New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.

Claim 4; Page 92; 119pp; English.

Compounds of the structure or formula S-(L) n-B, R-1-S-(L) n-R or S-(L) -R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4)); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-rP) (1-34); R-1 is the position of position of position of position of position of position of positions of positions of positions of positions of positions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases . binding R 1 is the (where

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The invention relates to a novel parathyroid hormone (PTH) peptide
CC (AAB01859) and parathyroid hormone-related peptide (PTHFP, AAB01860), and
CC biologically active derivatives thereof (AAB01857-B01858, AAB01861)
CC B01869). The peptides of the invention are at least 83 identical to the
CC generic peptide of the formula: X1-Val-Ser-Glu-Z2-Gln-Leu-X3-His-X4-X5-
CC Gly-Lys-X6 (AAB01857) where: X1 is Ser or Ala; X2 is 11e or Lys; X3 is
CC Met, Leu or Nle; X4 is Asn or Asp; X5 is Leu or Lys; X6 is His or Ser;
CC provided that the peptide is not PTHFP(1-14). The peptides of the
CC invention also encompass fragments of peptides of the invention
CC consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C-
CC terminal derivatives. FTH is a major regulator of calcium homeostasis,
CC and is necessary for the normal function of the gastrointestinal,
CC skeletal, neurological system, neuromuscular and cardiovascular systems.
CC it binds to both PTH-1 receptors on oscoblasts and renal tubular cells,
CC and to the recently identified PTH-2 receptor. PTH has a potent anabolic
CC effect on the skeleton, and mediates calcium reabsorption, enhances
CC phosphate clearance and vitamin D synthesis in the kidney. A homologous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               and disorders associated with decreased tether1 activity, increasing call a mammalian cell having PTH-1 receptors, or screening for a peptide non-peptide PTH (claimed). The new compound can be administered by inhalation unlike the large native PTH or PTHrP which avoids the need regular injections to treat osteoporosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Parathyroid hormone (PTH) peptides, PTH related peptides and the nucleic acids that encode them, useful for treating osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (GARD/)
(KRON/)
(POTT/)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-339693/29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gardella TJ,
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Search completed: May 18, 2004, 10:01:53 Job time : 46.75 secs

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                                                                                                                                                                                                           cc calls of protein hormones, the PTH-related proteins (PTHrP) mimic some of ct the renal and skeletal actions of PTH, and also bind to the PTH-1 CC receptor. They do not bind to the PTH-2 receptor. The peptides of the cinvention are either agonists of PTH-1 and PTH-2 receptors (AAB01858, AAB01861-B01866) or are PTH-1/PTH-2 receptor antagonists (AAB01857-CC B01869). PTH-1/PTH-2 receptor appoints are useful for the treatment of conditions characterised by a decrease in bone mass, such as consecoporosis. PTH-1/PTH-2 receptor antagonists are useful for treating medical disorders that arise from excessive or altered action of the PTH-1/PTH-2 receptor antagonists are useful for treating cc useful in the determination of races of bone formation, bone resorption cand/or bone remodelling in a patient. The peptides of the invention are minimised versions of PTH or PTH-P which are inexpensive to prepare by compentional synthetic chemistry, and can be delivered to a patient via non-parenteral routes. Sequences AAB01858 and AAB01861-B01866 represent cover.
                                                                                     Query Match
Best Local S
Matches 7
                                                                                                                                                                           Sequence 9 AA;
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                                                                                     Similarity 7; Conserv
AVSEIQLLH
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Listing first 45 summaries
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100.425 Million cell updates/sec
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            US-09-730-174A-5
US-09-730-174A-1
US-09-730-174A-1
US-09-730-174A-6
US-09-730-174A-6
US-09-730-174A-2
US-10-168-185-7
4 US-10-168-185-7
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6 US-10-168-185-7
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8 US-10-192-673-6
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Sequence 7, Appli
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Sequence 5, Application US/09730174A
Patent NO. US20020110871A1
GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and I
TITLE OF INVENTION: Selective Bind;
FILE REFERENCE: IMUNE-001A
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US-09-730-174A-5
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8-10-117-937-2	US-10-026-066-83	8-10-026-066-3	-10-094-699-4	-10-245-871-28	8-10-117-937-2	-094-699-5	0-253-2	-854-248-1	S-09-821-734-	-894-018-33	-10-374-035-1	-10-368-280-	-10-286-457-4	-10-286-457-29	-10-653-048-4	10-666-696-4	-10-645-761-4	0-651-723-4	S-10-632-388-	-10-609-217-	S-10-264-309-3	09-826-290-2	S-09-747-28	S-10-168-185-1	-10-168-185-	S-10-319-130-	-09-746-945	33-741-	US-10-192-673-8
equence 24	ence 83	equence 31	equence 45	equence 28	equence 24	ence 50	egue	quence 11	equence 4,	332	equence 12	quence 12	equence 46	quence 29	equence 41	equence 41	equence 41	equence 41	ence 41	equence 41	equence 36	nce 27,	quence 185, Ap	10e 11, Ap	ce 5, Appl	equence 16, App	nce 6, Appli	equence 61, App	Sequence 8, Appli

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; OTHER INFORMATION: Induces Formation of Antibodies and Isolates US-09-730-174A-3
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Patent No. US20020110871A1
GENERAL INFORMATION;
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having
TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid H
FILE REFERENCE: IMUNE-001A
CURRENT FILIGATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 3
LENGTH: 12
TYPE: PRT
CORGANISM: Artificial Sequence
and Peptide Antigens for Producing Antibodies having Binding Specificity to Bioactive Intact Parathyroid
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Pred. No. 0.00071;
; Mismatches 0;
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RESULT 4
US-09-730-174A-4
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US-09-730-174A-1
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                                                                                                               Sequence 4, Application US/09730174A

Patent No. US20020110871A1

GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having
TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid H
FILE REFERENCE: IMUNE-001A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Zahradnik, R.J.

TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid H FILE REFERENCE: IMUNE-001A CURRENT APPLICATION NUMBER: US/09/730,174A

CURRENT FILING DATE: 2000-12-05

NUMBER OF SEQ ID NOS: 12

SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1, Application US/09730174A Patent No. US20020110871A1 GENERAL INFORMATION:
   Query Match 93.2%;
Best Local Similarity 91.7%;
Matches 11; Conservative
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Best Local (
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CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 5
LENGTH: 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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Best Local Similarity
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                    OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                        FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
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                                                                                   Induces Formation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93.2%;
100.0%;
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 Score 55; DB 9; L
Pred. No. 0.0036;
0; Mismatches 1;
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Pred. No. 0.0024;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                    of Antibodies and Isolates
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                               Length 12,
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                                                                                    said Antibodies
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Sequence 1, Application US/10168185; Publication No. US20030175802A1; GENERAL INFORMATION:
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US-09-730-174A-2
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US-09-730-174A-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/09730174A
Patent No. US20020110871A1
GRMERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO 2
LENGTH: 11
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a
TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Hop
FILE REFERENCE: IMUNE-001A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
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Best Local S
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Best Local Similarity
Matches 10; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 6, Application US/09730174A
Patent No. US20020110871A1
APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinric
APPLICANT: Roth, Heinz-Jurgen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Antibodies and Peptide Antigens for TITLE OF INVENTION: Selective Binding Specificity to FILE REFERENCE: IMUNE-001A CURRENT APPLICATION NUMBER: US/09/730,174A CURRENT FILLING DATE: 2000-12-05 NUMBER OF SEQ ID NOS: 12
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TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
                                                                                                                                                                                                                                                                                                               Local Similarity
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                                                                                                                                                                                                                                                                                                               90.9%;
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                                                                                                                                                                                                                                                                                                               Score 51;
Pred. No.
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Pred. No. 0.012;
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                                                                                                                                                                                                                                                                                                                                Length 11;
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TITLE OF INVENTION:

Schmidt-Gayk, Heinrich

Heinz-Jurgen N: Method for Determining

Parathormone

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APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Uurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Method for Determining Parathormone
FILE REFERENCE: HLZ-004US
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TITLE OF INVENTION: Activity in a Human Sample FIL3 REFERENCE: HLZ-004US; CURRENT APPLICATION NUMBER: US/10/168,185; CURRENT FILING DATE: 2002-06-17; PRIOR APPLICATION NUMBER: PCT/EP00/12911; PRIOR FILING DATE: 2000-12-18; PRIOR APPLICATION NUMBER: DE 19961350; PRIOR APPLICATION NUMBER: DE 19961350; PRIOR APPLICATION NUMBER: DE 19961350; PRIOR FILING DATE: 1999-12-17; NUMBER OF SEQ ID NOS: 11; SOFTWARE: FastSEQ for Windows Version 4.0; SEQ ID NO 1
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-168-185-1
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US-10-168-185-7
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Best Local S
Matches 9
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                                             APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 9
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
mes 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Match 83.1%; So Local Similarity 100.0%; Fig. 10; Conservative 0;
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                                                                                                                                                                                                                                                                                       Sequence 6, Application US/10192673
Publication No. US20030166838A1
GENERAL INFORMATION:
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives
TITLE OF INVENTION: Brianthyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570002
CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
                                                                                                                                                                                      APPLICANT: Gardella, Thomas J. APPLICANT: Kronenberg, Henry APPLICANT: Potts, John T. APPLICANT: Juppner, Harald
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APPLICANT: Schmidt Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parachormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT APPLICATION UNMER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR TILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DC 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TYPE: PRT ; ORGANISM: Homo sapiens US-10-168-185-2
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 8, Application US/10168185 Publication No. US20030175802A1 GENERAL INFORMATION:
                                                                                                        Query Match
Best Local 9
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Best Local Similarity
Matches 9; Conserv
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PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Armbruster, Franz Paul APPLICANT: Missbichler, Albert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT FILING DATE:
                                                                                                                                                                                               LENGTH: 8
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENGTH:
                                                                                  'Match 69.5%; Score 41; DB 14; Local Similarity 100.0%; Pred. No. 1e+06; les 8; Conservative 0; Mismatches
                                         3 SEIQLMHN 10
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SEIQLMHN 8
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llarity 100.0%; Pred. No. 1e+06;
Conservative 0; Mismatches 0; Indels
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                                                                                                                                DB 14; Length 8;
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Sequence 7, Application US/10192673

Publication No. US20030166838A1

GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Jophner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Bioactive Peptide (PTH) and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)

FILE REFERENCE: 0609.4570002
CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: US. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 7
LENGTH: 9
CREANISM: Artificial Sequence
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US-10-192-673-7
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                                                                                                                                                                                                                 US-10-168-185-3
                                                                                                                                                                                                                                       RESULT 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-10-192-673-7
Sequence 3, Application US/10168185
Publication No. US20030175802A1
GENERAL INFORMATION:
APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Marches 8; Conserv
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Best Local
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PRIOR APPLICATION NUMBER: U.S.
PRIOR FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
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                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 77.
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88.9%;
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77.8%;
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Pred. No. 1e+0
2; Mismatches
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Pred. No. 1e+06;
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APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Bioactive Peptides and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609,4570002
CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR APPLICATION NUMBER: U.S. 50/105,530
PRIOR FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
LENGER: 0
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Best Local Similarity
7; Conserve
       ; Sequence 4, Application US/10168185
publication No. US20030175802A1
; GENERAL INFORMATION:
   APPLICANT: Armbruster, Franz Paul
   APPLICANT: Missbichler, Albert
   APPLICANT: Schmidt-Gayk, Heinrich
   APPLICANT: Roth, Heinz-Jurgen
   ITILE OF INVENTION: Method for Determining Parathormone
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US-10-168-185-4
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CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF 5BQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
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Publication No. US20030166838A1
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TYPE: PRT
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FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: DE 10961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTMARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 4
LENGTH: 7
TYPE: PRT
ORGANISM: Homo sapiens
US-10-168-185-4
Query Match
Best Local Similarity 100.0%; Pred. No. 1e+66;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0

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Db 1 SVSEIQL 7
Search completed: May 18, 2004, 10:20:59
Job time: 33.25 secs
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Sequence 31, Appl Sequence 2, Appli Sequence 2, Appli Sequence 6, Appli Sequence 3, Appli Sequence 3, Appli Sequence 4, Appli Sequence 4, Appli Sequence 14, Appli Sequence 14, Appli Sequence 15, Appli Sequence 16, Appli Sequence 16, Appli Sequence 17, Appli Sequence 18, Appli Sequence 18, Appli Sequence 18, Appli Sequence 18, Appli Sequence 14, Appli Sequence 17, Appli Sequence 13, Appli Sequence 13, Appli Sequence 13, Appli Sequence 13, Appli Sequence 12, Appli Sequence 14, Appli Se
                                                 ; FEATURE:
; NAME/KEY: MOD RES
; LOCATION: (8) 7
; CTHER INFORMATION: N
US-09-442-989-31
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APPLICANT: Mencel, James J.
PITIE OF INVENTION: PROCESS FOR THE PREPARAT
ITILE OF INVENTION: PEPTIDES
FILE REFERENCE: A31138-US
CURRENT APPLICATION NUMBER: US/09/442,989
CURRENT FILING DATE: 1999-11-18
EARLIER APPLICATION NUMBER: 60/081,897
EARLIER FILING DATE: 1998-04-15
NUMBER OF SEQ ID NOS: 46
SOFTWARB: PATENTIN Ver. 2.1
SEQ ID NO 31
LENGTH: 12
TYPE: PRT
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US-09-442-989-31
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Patent No. 6569993
GENERAL INFORMATION:
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LOCATION: (10)
CTHER INFORMATION:
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OTHER INFORMATION:
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US-09-287-221-6
US-09-287-221-6
US-08-140-137A-42
US-08-140-604-74
US-08-160-604-75
US-09-51-976-10
US-09-51-976-10
US-09-543-177
US-08-160-604-73
US-08-248-839-7
US-08-248-839-7
US-08-160-604-73
US-08-160-604-74
US-08-160-604-74
US-08-160-604-74
US-08-160-604-74
US-08-160-604-74
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42, Appl
42, Appl
271, App
74, Appl
75, Appl
10, Appl
10, Appl
117, Appl
117, Appl
127, Appl
127, Appl
180, Appl
23, Appl
24, Appl
25, Appl
27, App
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Result

Score

Query Match

Length

BB

Description

SUMMARIES

US-09-442-989-31
US-08-817-547A-1
US-08-817-547A-2
US-09-421-379-6
US-09-421-379-10
US-09-421-379-10
US-09-421-379-10
US-09-421-379-10
US-08-817-547A-1
US-08-817-547A-14
US-08-817-7109A-5
US-08-817-7109A-5
US-08-817-7109A-6
US-08-817-747A-17
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US-08-817-547A-15
US-08-817-547A-13
US-08-818-547A-13
US-08-818-547A-13
US-08-318-693-12

Minimum Maximum

DB BG

seq length: 0 seq length: 12

Total number of hits satisfying chosen parameters:

121490

389414 segs, 51625971 residues

Scoring table: Sequence: Title: Perfect score:

BLOSUM62

Gapop 10.0 ,

Gapext 0

US-09-730-174A-3 59

SVSEIQLMHNLG

12

Post-processing: Minimum Match Maximum Match Listing first

1 0% 1 100% 1 45 summaries

Database

Issued_Patents_AA:*

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6: /cgn2_6/ptodata/2/iaa/backfiles1.pep:*

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US-08-817-547A-2
                                                                                                                                                                          RESULT 3
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US-08-817-547A-1
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Sequence 2, Application US/08817547A
Patent No. 6030790
GENERAL INFORMATION:
APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Hock, Dieter
TITLE OF INVENTION: Peptides from the hPTH Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Appric Patent No. 6030790
                                                                                                                                                                                                                                                                                                                        Matches
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Best Local
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: PCT/EP95/
APPLICATION NUMBER: PCT/EP95/
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 0782
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 404-818-3700
TELEPAX: 404-818-3799
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: unkno
MOLECULE TYPE: po
HYPOTHETICAL: no
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/817,547A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Atlanta
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10; Conserv
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                                                                                                                                                                                                                                                                                                                   83.1%; Score 49; DB 3;
100.0%; Pred. No. 0.0057;
tive 0; Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07826-0007
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Pred. No. 0.
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                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Gardella, Thomas J.
APPLICANT: Cronenberg, Henry
APPLICANT: John T.
APPLICANT: Juppner, Harald
ITITE OF INVENTION: Bloactive Peptides and Peptide Derivatives of
ITITE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
ITITE OF INVENTION: Hormone-Related Peptide (PTHrP)
ITITE OF INVENTION: Hormone-Related Peptide (PTHrP)
ITITE OF INVENTION: HORMONE US/09/421,379
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: U.S. 60/105,530
EARLIER APPLICATION NUMBER: U.S. 50/105,530
  ; ORGANISM: Artificial Sequence; FEATURE:
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide
US-09-421-379-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                B
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Best Local Similarity
Matches 9; Conserv
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 07826-0007
TELECOMMUNICATION INFORMATION:
                                                                                                                           ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: unknown
MOLECULE TYPE: po
HYPOTHETICAL: no
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A
FILING DATE:
FILING DATE:
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MEDIUM TYPE: Floppy disk
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CITY: Atlanta
STATE: Georgia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 9 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
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Y: USA
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100.0%; Pred. No.
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3e+05;
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APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Boactive Peptide (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrp)
FILE REFERENCE: 0609,4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-09-421-379-7
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US-09-421-379-7
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Best Local :
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
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Patent No. 6495662
                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                    equence 3, Application US/08817547A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 9
                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                       STREET: 191 Pe
CITY: Atlanta
                                APPLICATION NUMBER: US/08/817,547A
                                                                                                                                                                      ZIP: 30303
                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                  ADDRESSEE:
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Y: USA
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191 Peachtree Street, 37th Floor
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77.8%;
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Pred. No. 3e+0
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 40; DB 4;
Pred. No. 3e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 4;
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APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Hotts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Boactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE REPERENCE: 0609.4570001
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: U.S. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
밁
                                                                                                                                         ; FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide
US-09-421-379-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; ANTI-SENSE: no
US-08-817-547A-3
                                                                   Best Loc
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 649566;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
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INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                     CENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 078
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HYPOTHETICAL:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T
                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: unl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: PCT/E
FILING DATE: 29 SEPT 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TO,
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                              1 SVSEIQLMH 9
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                                                                   Similarity 77.8
7; Conservative
SVSEHQLLH 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Application US/09421379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                59.3%; Score 35; DB nilarity 100.0%; Pred. No. 3e-Conservative 0; Mismatches
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                                                                                     55.9%;
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                                                                   Score 33; DB 4;
Pred. No. 3e+05;
1; Mismatches
                                                                                                        DB 4;
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                                                                                                      Length 9;
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                                                                 Gaps
                                                                   0;
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5460978-3
; PATENT NO. 5460978
; PATENTI MARTIN, THOMAS J.; MOSELEY, JANE M.; KEMP,
; BRUCE E.; WETTENHALL, RICHARD E.H.
; TITLE OF INVENTION: PROTEIN ACTIVE IN HUMORAL
; HYPERCALCEMIA OF MALIGNANCY-PTHRP

RESULT 8

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RESULT 9
US-08-817-547A-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 4, Application US/08817547A Patent No. 6030790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 54.2%;
Best Local Similarity 70.0%;
Matches 7; Conservative
                                                Query Match
Best Local
                                Matches
                                                                                                                                                                                                                                                                               PILING DATE:
PRIOR APPLICATION DATA:
PRIOR APPLICATION UNMBER: PCT/EP95/03757
PILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 07826-0007
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-818-3700
                                                                                                                                                                                                                                                    TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markue
TITLE OF INVENTION: Peptides
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 199,235
FILING DATE: 09-MAY-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/715,280
FILING DATE: 14-JUN-1991
                                                                                                               HYPOTHETICAL:
ANTI-SENSE: 1
                                                                                                                                          MOLECULE TYPE:
                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 4
                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: un
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 11
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1 SVSEIQL
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                              Similarity 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3: Jones & Askew, LLP
191 Peachtree Street, 37th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           USA
                              Conservative
                                                                                                                  no
                                                                                                                                                                 unknown
                                                                                                                                               peptide
                                                                                                                                                                                 unknown
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                                              50.8%; Score 30; DB 3; Length 7; 100.0%; Pred. No. 3e+05;
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                              0;
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Pred. No. 8.2;
1; Mismatches
                                Mismatches
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                              Gaps
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PRIOR APPLICATION DATA:

APPLICATION NUMBER: PCT/EP95/

FILING DATE: 29 SEPT 1994

ATTORNEY/AGENT INFORMATION:

NAME: FROST, ROSER T.

REGISTRATION NUMBER: 22,176

REFERENCE/DOCKET NUMBER: 0782

07826-0007

PCT/EP95/03757

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A
FILING DATE:

NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS: STREET: 191 Pe

from the hPTH Sequence

ADDRESSEE:

E: Jones & Askew, LLP 191 Peachtree Street, 37th Floor

COUNTRY:

USA

Georgia

30303

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APPLICANT: Gardella, Thomas J.
APPLICANT: Kroneaberg, Henry
APPLICANT: Notes, John T.
APPLICANT: Dotts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Normone-Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: U.S. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 8
LENGTH: 9
                                                                                                   Sequence 5, Application US/08817547A
Patent No. 6030790
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 8, Application US/09421379
Patent No. 6495662
                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides
                                                                                                                                                                                                                                                                                                                                            Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  μ
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6; Conservative
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                                                                                                                                                                                                                                                                                                                                         50.8%;
                                                                                                                                                                                                                                                                                                                                            Score 30; DB 4;
Pred. No. 3e+05;
                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                DB 4; Length 9;
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COUNTRY: USA
ZIP: 10016-2391
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/428,257A
FILING DATE: 07/05/95
CLASSIFICATION: 514
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 12
US-08-428-257A-14
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                                                                                                  RESULT 13
US-08-802-981-124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 14, Application US/08428257A Patent No. 5885808 GENERAL INFORMATION:
                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 5; Conserv
                                                                    Sequence 124, Application US/08802981 Patent No. 6037137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOC.
MOLECULE TYPE: pe
APPLICANT: Komoriya, Akira
APPLICANT: Packard, Beverly S.
TITLE OF INVENTION: Compositions for the Detection of Enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Jules E. Goldberg
STREET: 261 Madison Avenue
CITY: New York
STATE: NY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Spooner, Robert A.
APPLICANT: Epenetos, A.A.
TITLE OF INVENTION: Compounds to target cells
NUMBER OF SEQUENCES: 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL: I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION: TELEPHONE: 404-818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 100.0%; Pred. No. 3e+05; res 6; Conservative 0; Miamatahan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 6 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: unknown
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                                                                                                                                                                             3 TVSEVQL 9
                                                                                                                                                                                                           1 SYSEIQL 7
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                                                                                                                                                                                                                                             44.1%; Score 26; DB 2; Length 10; 71.4%; Pred. No. 94; rative 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 6;
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/177,109A
FILING DATE: 03-JAN-1994
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:

ADDRESSEE: Patrea L. Pabst STREET: 2800 One Atlantic Center STREET: 1201 West Peachtree Street CITY: Atlanta STATE: Georgia

```
NAME/KEY: Modified-site
; LOCATION: 3
; OTHER INFORMATION: /product= "Aib"
US-08-802-981-124
                                                                                                                                                    RESULT 14
US-08-177-109A-56
                                                                                  : Sequence 56, Application US/08177109A; Patent No. 5869615; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Hunter, Tom
REGISTRATION NUMBER: 38,498
REFERENCE/DOCKET NUMBER: 0168
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 124:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICANT: Dennis E. Hourcade and Teresa J. Oglesby TITLE OF INVENTION: MODIFIED COMPLEMENT PROTEASES NUMBER OF SEQUENCES: 62
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/08/802,981
FILING DATE: 20-FEB-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                               ration 42.4%; Score 25; DB 3; Length 11; Local Similarity 80.0%; Pred. No. 1.6e+02; es 4; Conservative 1; Mismatches n. Thank
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RESULT 15
US-08-687-706-56
; Sequence 56, Application US/08687706
; Patent No. 5928892
; GENERAL INFORMATION:
; APPLICANT: Dennis E. Hourcade an
                                                                                                                                                                                                         APPLICATION NUMBER: US/08/687,706
FILING DATE: 26-JUL-1996
CLASSIFICATION: 514
PRIOR APPLICATION NUMBER: US 08/177,109
FILING DATE: 03-JAN-1994
CLASSIFICATION NUMBER: US 08/177,109
FILING DATE: 03-JAN-1994
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Pabst, Patrea L.
REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: WU 107 DIV
REFERENCE/DOCKET NUMBER: WU 107 DIV
TELECOMMUNICATION INFORMATION:
TELEPHONE: (404) 873-8794
INFORMATION FOR SEQ ID NO: 56:
SEQUENCE CHARACTERISTICS:
LENGTH: Samino acidae
LENGTH: Samino acidae
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Query Match 40.7%; Score 24; DB 2; Length 5; Best Local Similarity 60.0%; Pred. No. 3e+05; Matches 3; Conservative 2; Mismatches 0; Indels
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Best Local Similarity 60.0%;
Matches 3; Conservative
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TELEPAX: (404) 873-8795
INFORMATION FOR SEQ ID NO: 56:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 30309-3450
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CONFUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Dennis E. Hourcade and Teresa J. Oglesby
TITLE OF INVENTION: MODIFIED COMPLEMENT PROTEASES
NUMBER OF SEQUENCES: 62
CORRESPONDENCE ADDRESS:
ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic Center
STREET: 1201 West Peachtree Street
CITY: Atlanta
STATE: Georgia
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TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
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REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: WU 107
TELECOMMUNICATION INFORMATION:
TELEPHONE: (404) 873-8794
                                                                                                                                                                               LENGTH:
TYPE: a
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Qy 8 MHNLG 12
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Db 1 LHNMG 5
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Search completed: May 18, 2004, 10:06:43 Job time : 12.75 secs

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Result
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Maximum Match 100%
Listing first 45 summaries
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                     of hits satisfying chosen parameters:
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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Match
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61
1 SVSEIQFMHNLG 12
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                           amine oxidase (cop
polyphosphate gluc
Ig heavy chain CDR
N-methylpurine DNA
alpha-2-macroglobu
hypothetical prote
lectin GNL1 alpha
rhodopsin homolog
type II site-speci
T-cell receptor al
T-cell receptor to
neuropeptide Antho
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kidney and bladder
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                                                                                                                                                                                                                                                                                                                     Description
                                                                                                                                                                                                                                                                                                                                               A;Gene: psbF
A;Genome: chloroplast
C;Superfamily: cytochrome b559 component
C;Keywords: chloroplast; photosynthesis;
                                                                                                                                                                                                                                                                                                                                                                                                                                    cytochrome b559 component psbF - pepper chloroplast (fragment)
C;Species: chloroplast Capsicum annuum (pepper)
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 11-Jun-1999
C;Accession: S28055
R;Kuntz, M.; Camara, B.; Weil, J.H.; Schantz, R.
Plant Mol. Biol. 20, 1185-1188, 1992
A;Title: The psbL gene from bell pepper (Capsicum annuum): plastid RNA editing also A;Reference number: S28055; MUID:93099270; PMID:1463853
A;Accession: S28055
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S28055
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A; Residues: 1-10 <KUN>
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ALIGNMENTS

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C;Accession: A58718
R;Stoffels, G.; Nissen-Meyer, J.; Gudmundsdottir, A.; Sletten, K.; Holo, H.; Nes, I.F.
Appl. Environ. Microbiol. 58, 1417-1422, 1992
A;Title: Purification and characterization of a new bacteriocin isolated from a Carnoback
A;Reference number: A58718; MUID:92321768; FMID:1622206
A;Accession: A58718.
                                                                     A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-7 <5TO>
C;Keywords: antibiotic; lanthionine
                                                                                                                                                                                                                                  C;Date: 23-Jan-1998 #sequence_revision 30-Jan-1998 #text_change 30-Jan-1998
C;Accession: A58718
                                                                                                                                                                                                                                                                     carnocin UI49 - Carnobacterium sp. (fragment)
C;Species: Carnobacterium sp.
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1 SISAMQFI 8
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                      Similarity
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   Conservative
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100.0%; 5
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                    29.5%; Score 18; 100.0%; Pred. No.
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Pred. No. 2.1e
3; Mismatches
     Mismatches
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2.1e+02;
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   2.8e+05;
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R;Yoshino, K.I.; Kajiura, H.; Nomura, K.; Takao, T.; Shimonishi, Y.; Kurita, Comp. Biochem. Physiol. B 94, 739-751, 1989
A;Title: A halogenated amino acid-containing sperm activating peptide and its otus nudus, Echinometra mathaei and Heterocentrotus mammillatus.
A;Reference number: A60527
                                                                                                                                         A;Resiques:
C;Keywords:
                                                                                                                                                                                                    R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rinehart Jr., Biochem. Biophys. Res. Commun. 124, 350-358, 1984
A;Title: Structures of two cockroach neuropeptides assigned by fast atom bombardment A;Reference number: A90118; MUID:85046530; PMID:6548628
A;Accession: A05169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Accession: S78420
R;Goldschmidt-Reisin, S.; Graack, H.R.
submitted to the Protein Sequence Database, February
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT
S78420
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                               80
                                                                                                                                                                 A; Molecule type: protein A; Residues: 1-8 <WIT>
                                                                                                                                                                                                                                                                                           neuropeptide M-I - American cockroach
C;Species: Periplaneta americana (American cockroach)
C;Date: 05-Jun-1987 #sequence_revision 05-Jun-1987 #t
C;Accession: A05169
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sperm-activating peptide (Tyr-2, Asn-3, Gly-5, Ile-9, Asp-10 SAP-I) - slate-pencil urchi
C;Species: Heterocentrotus mamillatus
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 18-Aug-2000
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A;Residues: 1-9 <GOL>
A;Note: the protein is designated as mitochondrial ribosomal protein
C;Keywords: mitochondrion; protein biosynthesis; ribosome
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A; Accession: S78420
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Feb-1998 #sequence_revision 13-Mar-1998 #text_change 21-Jul-2000
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EVNESPN 7
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Pred. No. 2.8e
1; Mismatches
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Pred. No. 2e+03;
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Pred. No. 2.8e+05;
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2.8e+05;
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C;Accession: S21205

R;Makiya, R.; Stigbrand, T.

Eur. J. Biochem. 205, 341-345, 1992

A;Title: Placental alkaline phosphatase has a binding site

A;Reference number: S21205; MUID:92209522; PMID:1555592

A;Accession: S21205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 7
221205
221205
19 heavy chain V region - human
C;Species: Homo sapiens (man)
C;Species: 22-Nov-1993 #sequence_revision 10-Nov-1995
C;Date: 22-Nov-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ig heavy chain C region (129) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1997 #sequence_revision 13-Mar-1998 #text_change
C;Accession: A37268
R;Ruff-Jamison, S.; Campos-Gonzalez, R.; Glenney Jr., J.R.
J. Biol. Chem. 266, 6607-6613, 1991
A;Title: Heavy and light chain variable region sequences and ant
A;Reference number: A38740; MUID:91177923; PMID:1706720
A;Accession: A37268
                                                                                                                             R;Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; Barra, J. Exp. Med. 176, 439-447, 1992
A;Title: H-2-restricted cytolytic T lymphocytes specific for HLA display T cell A;Reference number: S26512; MUID:92364546; PMID:1380061
A;Accession: PH1187
                                                                                                                                                                                                                 A;Residues: 1-12 <MAK>
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-10 <RUF:
                                                                      A; Molecule type:
A; Residues: 1-12
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                                                                                                           A; Status: preliminary
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Query Match
Best Local S
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               27.9%;
Score 17; DB Pred. No. 3.8e
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Pred. No. 3.1e+03;
1; Mismatches 4
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Pred. No. 3.8e+03;
3; Mismatches
                 DB 2;
3.8e+03
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SVSEIQF

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hypertrehalosemic hormone I - stick insect (Carausius morosus)
N,Alternate names: neuropeptide Cam-HrTH-I
N,Contains: hypertrehalosemic factor II
C.Species: Carausius morosus
C.Date: 30-Sep-1993 #sequence revision 30-Sep-1993 #text_change
C.Accession: JC1416; S07157
R.Gaede, G.; Kellner, R.; Rinehart, K.L.; Proefke, M.L.
Biochem. Biophys. Res. Commun. 189, 1303-1309, 1992
A,Title: A tryptophan-substituted member of the AKH/RPCH family
A,Reference number: JC1416; MUID:93129188; PMID:1482345
A,Accession: JC1416
         A,Molecule type: protein
A,Residues: 1-10 <GRE1>
A,Residues: 1-10 <GRE1>
R;Gaede, G.; Rinehart Jr., K.L.
Biol. Chem. Hoppe-Seyler 368, 67-75, 1987
A,Title: Primary structure of the hypertrehalosaemic factor
A,Reference number: S07157; MUID:87157103; PMID:3828078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Binette, J.P.; Binette, M.B. submitted to the Protein Sequence Data A;Description: The proteins of kidney
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Friedman, D.J.; Umeda, P.K.; Sinha, A.M.; Hsu, H. Froc. Natl. Acad. Sci. U.S.A. 81, 3044-3048, 1984
A;Title: Characterization of genomic clones specifying rabbit alpha-A;Reference number: 146868; MUID:84221901; PMID:6328491
A;Accession: I46868
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(;Species: Oryotolagus cuniculus (domestic rabbit)
C;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change 05-Nov-1999
C;Accession: 146868
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A; Accession: G58502
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C;Species: unidentified bacterium
C;Date: 07-Feb-1997 #sequence_revision 07-Feb-1997 #text_change 10-Jul-1998
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Pred. No. 2.8e+05;
4; Mismatches 1
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Pred. No. 2.8e+05;
1; Mismatches 1
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iney and gallbladder stones
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N,Alternate names: C:
C;Species: Extatosom:
C;Date: 30-Jun-1992:
C;Accession: S09138
R;Gaede, G.; Rinehar
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Biol. Chem. Hoppe-Seyler 371, 345-354, 1990
A;Title: Primary structures of hypertrehalosaemic neuropeptides
entalis and of the stick insect Extatosoma tiarratum assigned by
A;Reference number: S08995; MUID:90253659; PMID:2340112
A;Accession: S09138
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A;Residues: 'Z',2-10 'GARE's
C;Comment: Hypertrehalosemic factor II lacks the tryptophan modification.
C;Comment: This peptide raises hemolymph levels of trehalose in the cockroach Periplanet
C;Superfamily: adipokinetic hormone
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; glycoprotein; hormone; neuropeptide; pyroglutamic aci
C;Keywords: endiated carboxyl end; Gln) #status experimental
F;1/Modified site: pyrolidone carboxylic acid (Gln) #status experimental
F;8/Binding site: carbohydrate (Trp) (covalent) #status experimental
F;10/Modified site: amidated carboxyl end (Thr) #status experimental
                                                                                A; Molecule type: protein A; Molecule type: protein A; Residues: 1-10 <GAE> A; Residues: 1-20 <GAE> A; Rote: the amino-terminal residue forms pyrrolidone carboxylic acid; therefore, A; Note: the amino-terminal residue formone C; Supperfamily: adipokinetic hormone O; Keywords: amidated carboxyl end; hormone; neuropeptide; pyroglutamic acid C; Keywords: amidated carboxylic acid (Gln) #status experimental F; 10/Modified site: amidated carboxyl end (Thr) #status experimental
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                                                                                                                                                                                                           Biochem. Biophys. Res. Commun. 159, 726-733, 1989
A;Title: Primary structure of a pyrroloquinoline quinone
A;Reference number: A32428; MUID:89193662; PMID:2539124
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33.3%; Pred. No. 4.8e+03;
7. Mismatches 4;
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                    26.2%;
  Score 16; DB Pred. No. 5.3e 2; Mismatches
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Pred. No. 4.8e+03;
2; Mismatches 4
                                           DB 2;
                      .3e+03
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SEIQFMHN 10

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ig heavy chain CDR3 region (clone 1-118B) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep_1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0231
R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
U. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
A;Accession: PT0231
A;Molecule type: DNA
A,Recidues: 1-9 <YAM-
A;Residues: 1-9 <YAM-
A;Residues: 1-9 <YAM-
A;Residues: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Status: preliminary
A;Molecule type: protein
A;Residus: 1-7 cHI's
C;Keywords: phosphotransferase
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Search completed: May 18, 2004, 10:05:40 Job time : 10.25 secs
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polyodon sp
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litoria rub
bothrops ja

eisenia foe periplaneta bacteroides

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Minimum DB
Maximum DB
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Listing first
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re greater than or equal to the score of the result being printed,
is derived by analysis of the total score distribution.
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298
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Gapop 10.0 ,
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T2PI PROVU
FLRN ANTEL
ALL1 CYDPO
TY51 LITMU
ALL1 CYDPO
HTF1 PERAM
FAR8 MACRS
UF02 MOUSE
ALL1 CARMA
TKNC RANCA
TKNC PIG
ACST
HCY1 CARMA
TKNC PIG
HCY1 CARMA
LMT1 LOCMI
PORD METIM
RS19 PSEPU
HTF TENMO
LCK6 LEUMA
CONG CONGE
MOSH CLYJA
                                                                                                                                                                                                                                                                                                                                                                                                                LANC_CARUI
HTF1 ROMMI
ALL6_CYDPO
HTF2_CARMO
E103_LITRU
FUSS_FUSSO
EREL_PHODV
EFG_CLOPA
NUHM_CANFA
T2P1_PROVU
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   p49820 canis famil
p3103 proteus vul
p58707 anthopleura
p82158 cydia pomon
p82056 litoria rub
p81821 carcinus ma
p82152 cydia pomon
p0458 periplaneta
p83281 macrobrachi
p38640 mus musculu
p81822 carcinus ma
p22690 rana catesb
p01292 sus scrofa
p83146 bacteroides
p83146 bacteroides
p83176 carcinus ma
p22691 tomato big
p80903 methanobact
p25419 tenebrio mo
p21143 leucophaea
p1988 leucophaea
p19882 clypeaster
                                                                                                                                                                                                                                                                                                                                                                                                                                               Q03367 capsicum an p36960 carnobacter P18110 romalea mic P82157 cydia pomon P11385 carausius m p8209 litoria rub P81010 fusarium so P35946 phocine dis P81350 clostridium
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                                                                                                                                                  RESULT 1
PREF CAPAN
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ID PREF CAPAN
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OF O1-JUN
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Q03367;
01-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN=cv. Lamuyo; TISSUE=Fruit, and Leaf;

MEDLINE=93099270; pubMed=1463853;

Kuntz M., Camara B., Weil J.-H., Schantz R.;

"The psbL gene from bell pepper (Capsicum annuum): plastid RNA editing also occurs in non-photosynthetic chromoplasts.";

plant Mol. Biol. 20:1185-1188(1922).

-i- FUNCTION: This b-type cytochrome is tightly associated with the reaction center of photosystem II and possibly is part of the water-oxidation complex.

-i- SUBUNIT: Heterodimer of an alpha subunit and a beta subunit.

-i- SUBUNIT: Belongs to the psb2 / psbF family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Capsicum annuum (Bell pepper).
Chloroplast.
Endaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Eukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; asterids;
Iamiids; Solanales; Solanaceae; Capsicum.
                                                                                                                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                         NON TER
                                                                                                                                                                                                                          HAMAP, MF 00643; -; 1.
InterPro; IPRO6216; Cyt_b559.
PROSITE; PS00537; CYTOCHROME B559;
Chloroplast; Photosystem II; Heme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-JUN-1994 (Rel. 29, Created)
01-JUN-1994 (Rel. 29, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Cytochrome b559 beta subunit (PSII reaction c
                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                    EMBL; X65570; CAA46539.1; -. PIR; S28055; S28055.
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PPKI PERAM
ASL2_BACSE
CORZ_PERAM
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PVKI_PERAM
Q2OA_COWNTS
UKA2_HUMAN
UR2_FOLSP
REZI_LITRU
RE31_LITRU
ANG2_BOTJA
                                                                              Score 23; DB
Pred. No. 1.2e
3; Mismatches
                                                                                                                                                      BY SIMILARITY.
LUMENAL (POTENTIAL).
817D0F59D6D69DC5 CRC64;
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RESULT 2
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NON TER
SEQUENCE 7
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01-JUN-1994
16-OCT-2001
                                                                                                                                                                                                                                                                                             MEDINE-89145002; PubMed-3226948;
Gaede G., Hilbich C., Beyreuther K., Rineha "Sequence analyses of two neuropeptides of two heluber grasshopper, Romalea microptera. Peptides 9:681-688 (1988)
                                                                                                                      Neuropeptide; Amidation; Flight;
MOD_RES 1 1 PYRI
MOD_RES 10 10 AMII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lantibiotic car
Carnobacterium
                                                                                                      SEQÜENCE
                                                                                                                                                                               PROSITE; PS00256; AKH; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Neoptera; Orthopteroidea; Orthop
Acridoidea; Romaleidae; Romalea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Romalea microptera (Lubber grasshopper).
Eukaryota, Metazoa, Arthropoda, Hexapoda, Insecta, Pterygota,
Neoptera, Orthopteroidea, Orthoptera, Caelifera, Acridomorpha,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-NOV-1990 (Rel. 16, Created)
01-FEB-1994 (Rel. 28, Last sequence un
10-OCT-2003 (Rel. 42, Last annotation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=92321768;
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                                                                                                                                                                                                   InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                             SSUE=Corpora
                                                                                                                                                                                               FUNCTION: Hypertrehalosaemic factors are neuropeptides that elevate the level of trehalose in the hemolymph (trehalose major carbohydrate in the hemolymph of insects). SUBCELLULAR LOCATION: Secreted. SIMILARITY: Belongs to the AKH / HRTH / RPCH family. erPro; IPR002047; AKH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FUNCTION: Lanthionine-containing Active on Gram-positive bacteria. Ibiotic; Bacteriocin; Lantibiotic. TER 7
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                                                                                                      10 AA;
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carnocin UI49 (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                cardiaca;
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en-Meyer J., Gudmundsdottir A.,
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1163 MW;
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Pred. No.
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Pred. No. 1.1e+03;
2; Mismatches ;
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056236745771A9C4 CRC64;
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                                                                                                                                                                                                                                                                                                                                lart K.L. Jr.;
the AKH/RPCH-family
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                                                             Length 10,
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HTF2_CARMO
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f 01-VUL-1989 (Rel. 11, Created)
f 01-FEB-1994 (Rel. 28, Last sequence update)
f 00-FCT-2003 (Rel. 42, Last annotation update)
f 10-CCT-2003 (Rel. 42, Last annotation update)
f Hypertrehalosaemic factor II (HTF-II) (HRTH-II)
f Hypertrehalosaemic factor II (HTF-II) (HRTH-II)
f Carausius morosus (Indian stick insect), and
forestoama tiaratum (Stick insect).
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Best Local S
Matches 2
SEQUENCE.

SPECIES-E. tiaratum; TISSUE=Corpora cardiaca;

SPECIES-E. tiaratum; TISSUE=Corpora cardiaca;

MEDLINE=90253659; PubMed=2340112;

Gaede G., Rinehart K.L. Jr.;

Gaede G., Rinehart K.L. Jr.;

"Primary structures of hypertrehalosaemic neuropeptides isolated the corpora cardiaca of the cortoraches Leucophaea maderae, the corpora cardiaca of the corpora cardiaca.
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30-MAY-2000
30-MAY-2000
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MEDLINE=98054539; PubMed=9392829;

Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., W
Davey M., Bast P.D., Thorpe A.;

"Lepidopteran peptides of the allatostatin superfam
Peptides 18:1301-1309(1997)

-!- SIMILARITY: Belongs to the allatostatin family.
                                                                                                                                                                           Gaede G., Rinehart K.L. Jr.;
"Primary structure of the hypertrehalosaemic factor II
"Primary structure of the Indian stick insect, Carausius
determined by fast atom bombardment mass spectrometry."
Biol. Chem. Hoppe-Seyler 368:67-75(1987).
                                                                                                                                                                                                                                                      SPECIES=C.morosus; TISSUE=Corpora MEDLINE=87157103; PubMed=3828078; Gaede G., Rinehart K.L. Jr.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cydia pomonella (Codling moth).
Eukaryota, Metazoa; Arthropoda; Hexapoda; Insecta; Pteryg
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
                                                                                                                                                                                                                                                                                                                                                Heteronemiidae; Carausius.
NCBI_TaxID=7022, 7024;
                                                                                                                                                                                                                                                                                                                                                                                       Extatosoma tiaratum (Stick insect).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroidea; Phasmatodea; Euphasmida;
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P82157;
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0B2879C45B573767 CRC64;
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Pred. No. 1.4e+05
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(HRTH-II)
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Length

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Winstanley

D.

Pterygota;

Indels

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Gaps

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(Hypertrehalosaemic

Pterygota; Phasmatoidea;

morosus from the

orientalis

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MEDLINE-93129188; PubMed=1483345;

Gaede G., Kellner R., Rinehart K.L. Jr., Proefke M.L.;

Gaede G., Kellner R., Rinehart K.L. Jr., Proefke M.L.;

"A tryptophan-substituted member of the AKH/RPCH family isolated
a stick insect corpus cardiacum.";

Biochem. Biophys. Res. Commun. 189:1303-1309(1992).

-!- FUNCTION: Hypertrehalosaemic factors are neuropeptides that
-!- elevate the level of trehalose in the hemolymph (trehalose i
major carbohydrate in the hemolymph of insects).

-!- SUBCILIULAR LOCATION: Secreted.

-!- SUBCILIULAR LOCATION: Secreted.

-!- SUBCILIULAR LOCATION: Secreted.

-!- SUMILARITY: Belongs to the AKH / HRTH / RPCH family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       28-FEB-2003
28-FEB-2003
10-OCT-2003
                                                                                                                                                                                          Amphibian
MOD_RES
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EI03_I
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SPECIES=C.morosus; TISSUE=Corpora cardiaca;
                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                  Aust
                                                                                                                                                                                                                                                                                                                     TISSUE=Skin secretion;
Wabnitz P.A., Bowie J.H., Tyler M.J.,
"Peptides from the skin glands of the
Litori electrica. Comparison with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=104895;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata;
Amphibia; Batrachia; Anura; Ne
Pelodryadinae; Litoria.
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PROSITE; PS00256; AKH; 1.
Neuropeptide; Amidation; Glycoprotein; Pyrrolidone carboxylic acid.
Neuropeptide; Amidation; Glycoprotein; Pyrrolidone carboxylic acid.
MOD_RES 1 pyrroliDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                litoria rubella (Desert tree frog)
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m bombardment
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TISSUE SPECIFICITY: Skin.
Thibian defense peptide; Amidatio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JC1416; JC1416.
S09138; S09138.
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3; Conserv
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HVA
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Hoppe-Seyler 371:345-354(1990).
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(Rel.
(Rel.
                                                                                   Conservative
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41,
42,
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AMIDATION
                                                                               Score 15; DB 1;
Pred. No. 1.4e+05;
1; Mismatches (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Craniata; Vertebrata;
Neobatrachia; Hyloidea;
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Pred. No. 2.7e+03;
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Australian buzzing tree frog
skin peptides from Litoria
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family isolated
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Hylidae;
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RESULT

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RRPI_PHODV
P35946;
01-JUN-1994
01-JUN-1994
10-OCT-2003
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15-JUL-1998 (Rel. 36, Last sequence up
10-OCT-2003 (Rel. 42, Last annotation
Allergen Fus s I3596* (Fragment).
Fusarium solani (subsp. pisi) (Nectria
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MEDLINE=9228877; PubMed=1588321;

Curran M.D., O'Loan D., Kennedy S., Rima B.K.;

Curran M.D., O'Loan D., Kennedy S., Rima B.K.;

"Molecular characterization of phocine distemper virus: gene order and sequence of the gene encoding the attachment (H) protein.";

J. Gen. Virol. 73:1189-1194 (1992).

-i- FUNCTION: Probable component of the active polymerase. It may function in mRNA synthesis, capping, methylation and poly(A) synthesis of newly synthesized viral mRNAs, RNA editing of the gene transcript, and protein kinase activity.

-i- CAPALYTIC ACTIVITY: N nucleoside triphosphate = N diphosphate
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fr. protein) (Fragment).
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NON_TER
                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstatton the European Bioinformatics Institute. There are no restrictions on its use by mon-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=IARI 3596; TI:
Verma J., Gangal S.V
Submitted (JUL-1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hypocreomycetidae; Hypocreales; Nectriaceae; Nectria NCBI_TaxID=70791;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fusarium solani (subsp. pisi) (Nectria haemato
Eukaryota; Fungi; Ascomycota; Pezizomycotina;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P81010;
15-JUL-1998
Transferase; RNA-directed RNA
NON TER 10 10
SEQUENCE 10 AA; 1105 MW; 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                               EMBL; D10371; BAA01208.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=11240;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Phocine distemper virus (PDV).
Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -!- ALLERGEN: Causes an allergic reaction in human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                     SIMILARITY: Belongs to the paramyxoviruses L protein family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FUSSO
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HNV 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HNL 11
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(Rel. 29,
(Rel. 42,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TISSUE=Mycelium;
5.V.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          subunit (EC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 24.6%;
66.7%;
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Last annotation updat
ubunit (EC 2.7.7.48) (
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Pred. No. 1.4e
1; Mismatches
                                                    polymerase
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.48) (Large
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RESULT 10
NUHM CANFA
ID NUHM C
AC P49820
DT 01-OCT
DT 15-JUL
DT NADH-U
DE (BC 1.
GN NDUFV2
OS CANIS
OC EUKARY
OC MAMMAJ
OC NCBI []
RN [1]
RN [1]
RX MEDLIJ
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RA DUNN |
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RT Gos h
RL Elect
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EFG_CLOPA
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Best Local 9
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Best Local 9
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P81350;
15-JUL-1998
15-JUL-1998
28-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Flengsrud R., Skjeldal L.;
"Two-dimensional gel electrophoresis separation and N-terminal sequence analysis of proteins from Clostridium pasteurianum W5.";
Electrophoresis 19:802-806(1998).
-!- FUNCTION: This protein promotes the GTP-dependent translocation the nascent protein chain from the A-site to the P-site of the
                                                                                                                                                                                                  P49820;
01-OCT-1996
TISSUE-Heart; PubMed=9504812; PubMedINE-98163340; PubMed=9504812; Punn M.J., Corbett J.M., Wheeler C.H.; "HSC-2DPAGE and the two-dimensional gedge heart proteins."; Electrophoresis 18:2795-2802(1997).
                                                                                                                                                                                                                                                                                                                                                                                       EF-G/EF-2 subfamily.

InterPro; IPR000795; EF GTPbind.
PROSITE; PS00301; EFACTOR GTP; PARTIAL.
Elongation factor; Protein biosynthesis;
NON_TER 11 11
                                                                                                                                                       01-OCT-1996 (Rel. 34, Created)
15-JUL-1998 (Rel. 36, Last sequence up
10-OCT-2003 (Rel. 42, Last annotation
NADH-ubiquinone oxidoreductase 24 kDa
(EC 1.6.99.3) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                               - -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE=98291870; PubMed=9629918; Flengsrud R., Skjeldal L.;
                                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN-W5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=1501;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bacteria; Firmicutes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Clostridium
                                                                             SEQUENCE
                                                                                                NCBI_TaxID=9615;
                                                                                                                                   Canis familiaris
                                                                                                                                             NDUFV2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Elongation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           lostridium
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SIMILARITY: Belongs to
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                                                                                                                                                                                                                                                                                    LEKFONIG
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98 (Rel. 36, Last s
03 (Rel. 41, Last a
n factor G (EF-G)
                                                                                                                       Metazoa;
                                                                                                                                                                                                                                                                                                                                                                            11 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               pasteurianum.
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                                                                                                                                                                                                                            STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Belongs to the
                                                                                                      (Dog).
.oa; Chordata; Craniata; Ver
.oa Carmivora; Fissipedia;
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                                                                                                                                                                                                                                                                                                                                                                            1337 MW;
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25.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cytoplasmic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (CP 5) (Fragment)
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                                                                                                                                                                                                                                                                                                                               3; Mismatches
                                                                                                                                                                                                                                                                                                                                                                             412E71F1D9C33B17 CRC64;
                                                                                                                                                                                                                                                                                                                                                     Score 15;
                                                                                                                                                                                                                            PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                             GTP-binding
                                                                                                                                                                                                                                                                                                                                           No. 4.7e+03;
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                                                                                                                                                                                          update)
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subunit
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                                                                                                              Vertebrata;
ia; Canidae;
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                                 electrophoresis database
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                                                                                                                                                                     (E)
                                                                                                                                                                                                                                                                                                                                                     Length 11;
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                                                                                                               Euteleostomi;
Canis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              translocation
P-site of the
                                                                                                                                                                                                                                                                                                                                                                                                                                               factor family
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Query Match Best Local S Matches 3

24.6%;
Similarity 60.0%;
3; Conservative

Pred. No. 4.76 1; Mismatches Score 15; Pred. No.

4.7e+03; DB 1;

Length 11;

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Gaps

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1300 MW;

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CRC64;

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Query Match
Best Local
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Restriction
NON TER
SEQUENCE 1
                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the EN the European Bioinformatics Institute. There are no restructed by non-profit institutions as long as its content modified and this statement is not removed. Usage by and entities requires a license agreement (See http://www.isb-sentities.requires and the EN the EN
                                                                                                                                                                                                                                                                                                                                                         EMBL; L04163; AAA25660.1;
PIR; S35490; S35490.
REBASE; 1541; PvuI.
                                                                                                                                                                      or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WEDLINE=93087186; PubMed=1454536; Smith M.D., Longo M., Gerard G.F., Chatterjee D.K.; Smith M.D., Longo M., Gerard G.F., Chatterjee D.K.; "Cloming and characterization of genes for the PvuI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales;
Enterobacteriaceae; Proteus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUL-1993 (Rel 26, Creat
01-JUL-1993 (Rel 26, Last
10-OCT-2003 (Rel 42, Last
Type II restriction enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oxidoreductase; NAD; Ubiquinone; Mitochondrion; Iron-sulfur; Iron; 2Fe-25.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -!- SIMILARITY: Belongs
HSC-2DPAGE; P49820; DOG.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. STRAIN=ATCC 13315;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=585
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PROSITE; PS01099; COMPLEX1_24K; PARTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Proteus vulgaris.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PVUIR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FUNCTION: TRANSFER OF ELECTRONS FROM NADH TO THE RESPIRATORY CHAIM. THE IMMEDIATE ELECTRON ACCEPTOR FOR THE ENZYME IS BELL TO BE UBIQUINONE. COMPONENT OF THE FLAVOPROTEIN-SULFUR (FP) FRACKENT OF THE ENZYME.

CATALYTIC ACTIVITY: NADH + ubiquinon = NAD(+) + ubiquinol.

CATALYTIC ACTIVITY: NADH + acceptor = NAD(+) + reduced accept COFACTOR: Binds 1 2Fe-2S cluster (Potential).

SUBUNIT: Mammalian complex I is composed of 45 different subsubcellular LOCATION: Matrix and cytoplasmic side of the mitochondrial inner membrane.

SIMILARITY: Belongs to the complex I 24 kDa subunit family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROVU
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                                                  system;
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                                                     Hydrolase; Nuclease;
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annotation update)
PvuI (EC 3.1.21.4)
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Pred. No. 4.7e+03;
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(See http://www.isb-sib.
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                                                                                                                                                                                                                                                                                                              EMBL outstation
                                                                                                                                                                                                                                                                                                                                 a collaboration
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Best Local S
Matches 2
Query Match
Best Local Similarity
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P82158;
30-MAY-2000 (Rel. 3
30-MAY-2000 (Rel. 3
30-MAY-2000 (Rel. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FLRN_ANTEL STANDARD
P58707;
28-FEB-2003 (Rel. 41, C
-28-FEB-2003 (Rel. 41, I
10-CCT-2003 (Rel. 42, I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Neuropeptide;
MOD_RES
MOD_RES
                                                                                 MOD RES
SEQUENCE
                                                                                                              MEDLINE=98054539; PubMed=9392829;
Duve H., Johnsen A.H., Masstro J.-L., Scott.
Davey M., East P.D., Thorpe A.;
"Lepidopteran peptides of the allatostatin speptides 18:1301-1309(1997).
-- SIMILARITY: Belongs to the allatostatin Neuropeptide; Amidation.
MOD_RES 7 AMIDATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CYDPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anthopleura elegantissima (Sea anemone).
Eukaryota; Metazoa; Cnidaria; Anthozoa;
Nynantheae; Actiniidae; Anthopleura.
NCBI_TaxID=6110;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Cydia pomonella (Codling moth).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Grimmelikhuijzen C.J.P., Rinehart K.L. Jr., Jacob E., Graff D., Reinscheid R.K., Nothacker H.-P., Staley A.L.; "Isolation of L-3-phenyllactyl-Leu-Arg-Asn-NH2 (Antho-RNamide),
                                                                                                                                                                                                                                                                                                                                SEQUENCE.
TISSUE=Larva;
                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=82600;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Neuron specific.
-!- MASS SPECTROMETRY: MW=549.3; METHOD=FAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     group."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                anemone neuropeptide containing an unusual amino-terminal blocking
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE, AND MASS SPECTROMETRY. MEDLINE=90319122; PubMed=1973541;
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                                                                                                           leuropeptide;
fod_RES
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nilarity 50.0%;
Conservative
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. 39, Last seqn
. 39, Last anno
                                                                                    873 MW;
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  23.0%;
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Last annotation updat
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Last annotation updat
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                                                                                                                                                                                                                    the allatostatin superfamily.";
                                                                                 AMIDATION.
672879CABB569350 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          64540729A0000000 CRC64;
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Pred. No. 1.4e+05;
1; Mismatches 1
  Score
Pred.
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     14;
No.
                                                                                                                                                                                                                                                                           Scott A.G., Winstanley D.,
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     DB 1;
1.4e+05;
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                                                                                                                                                                     family
                                Length 7;
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Best Local S
Matches 2
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P81821;
30-MAY-2000
30-MAY-2000
30-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOD RES
MOD RES
SEQUENCE
  "Isolation and identification of multiple neuropeptides allatostatin superfamily in the shore crab Carcinus maes Eur. J. Biochem. 250:727-734 (1997).

-I- FUNCTION: May act as a neurotransmitter or neuromodu-
-I- SIMILARITY: Belongs to the allatostatin family.

Neuropeptide; Amidation; Multigene family.
                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyu
Eubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                                                                                                                          TISSUE=Cerebral ganglion, and Thoracic ganglion, MEDLINE=98121193; PubMed=9461295; Duve H., Johnsen A.H., Maestro J.-L., Scott A.G.
                                                                                                                                                                                                                                                                                                                                                                                                                                           Carcinustatin 18.
Carcinus maenas (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LITRU
                                                                                                                                                                        Thorpe A.;
                                                                                                                                                                                                                                                                                                                                     CBI_TaxID=6759;
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STANDARD;

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39, Created)
39, Last sequence 39, Last annotations

Last sequence update)

update)

(Common shore crab) (Green crab)

Brachyura;

Scott A.G.,

Jaros P.P.,

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Matches
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P82065;
28-FEB-2003 (Rel.
28-FEB-2003 (Rel.
10-OCT-2003 (Rel.
                                                                                                                                                                                                          Tyler M.J., Wallace J.C.;

"The structure of new peptides from the Australin red tree frog
'Litoria rubella'. The skin peptide profile as a probe for the study
of evolutionary trends of amphibians.";
Aust. J. Chem. 49:955-963(1996).

-!- FUNCTION: May act as a neuromodulator or neurotransmitter.

-!- SUBCELLULAR LOCATION: Secreted.

-!- TISSUE SPECIFICITY: Expressed by the skin dorsal glands.

-!- TISSUE SPECIFICITY: Expressed by the skin dorsal glands.

-!- MASS SPECIFONETRY: MW=965; METHOD=FAB.

Amphibian defense peptide; Amidation; Neuropeptide;

Pyrrolidone carboxylic acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tryptophyllin 5.1.
Litoria rubella (Desert tree frog).
Litoria rubella (Desert tree frog).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata;
Eukaryota, Metazoa; Chordata; Craniata; Hyloidea;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea;
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                             TISSUE=Skin secretion;
Steinborner S.T., Wabnitz P.A., Waugh R.J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=104895;
                                   4 EIQFMH 9
                                                                        Similarity
2; Conserv
QIPWFH 6
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                                                                                                                                                      7 AA;
                                                                          Conservative
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41,
42,
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                                                                                               23.0%;
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Last sequence update)
Last annotation updat
                                                                                                                                                         MW;
                                                                                                                                                      AMIDATION.
7401E9D3676046B0
                                                                            Score 14; DB
Pred. No. 1.4e
2; Mismatches
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                                                                                                               DB 1;
                                                                      .4e+05;
2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Bowie J.H.,
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                                                                                                                 Length
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                                                                            Gaps
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FT MOD RES 8 AMIDATION (POTENTIAL).

SQ SEQUENCE 8 AA, 919 MW; C82879D5AB569AB5 CRC64;

Query Match 23.0%; Score 14; DB 1; Length 8;

Best Local Similarity 40.0%; Pred. No. 1.4e+05;

Matches 2; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 8 MHNLG 12 | : : | |
Db 3 MYSFG 7

Search completed: May 18, 2004, 10:02:34

Job time: 7.75 secs

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Minimum
Maximum
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No.
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Maximum Match 100%
Listing first 45 s
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    of hits satisfying chosen parameters:
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1: sp_archea:*
2: sp_bacteria:*
3: sp_fungi:*
4: sp_inman:*
6: sp_mammal:*
6: sp_mammal:*
7: sp_mhc:*
9: sp_blage:*
10: sp_blant:*
11: sp_virus:*
12: sp_virus:*
13: sp_virus:*
14: sp_unclassified:*
15: sp_bacteriap:*
16: sp_bacteriap:*
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Match
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125.164 Million cell updates/sec
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Copyright (c) 1993 - 2004 Compugen Ltd.
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                                       Opps71 agkistrodon
Opt178 canis famil
Op2766 canine dist
071066 canine dist
P82081 limnodynast
O13591 saccharomyc
O15342 homo sapien
O8mesO anoda crist
O85631 avian carri
O23876 dictyosteli
O23874 paramecium
O28742 oryctolagus
O853120 gallus gall
Q05403 saccharomyc
Q40659 oryza sativ
P82082 limnodynast
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 RESULT
Q9TT78
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Best Local S
Matches 4
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NON TER 11
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11
SEQUENCE 11 AA; 1209 MW; 7CA02D1D41E8772B CRC64;
                                                                                                                                                                                                                                                                                                                                                                    Q9PS71 PRELIMINARY; PRT; 11 AA.
Q9PS71;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-UN-2000 (TrEMBLrel. 21, Last annotation update)
01-UN-2002 (TrEMBLrel. 21, Last annotation update)
Fibrinolytic metalloproteinase (Fragment).
Agkistrodon contortrix.
Bukaryota; Metazoa; Chordata, Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea;
Viperidae; Crotalinae; Agkistrodon.
WCBI TaxID=8720;
 Q9TT78;
Q9TT78;
Q1-MAY-2000 (TrEMBLrel. 13,
Q1-MAY-2000 (TrEMBLrel. 13,
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                                                                                                                                                                      39.3%; Score 24; DB 13; 100.0%; Pred. No. 5.9e+02; ative 0; Mismatches 0;
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Q34994
Q34993
Q43960
Q153960
Q158396
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Q44377
Q44468
Q8RKU3
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Q4446 aeromonas v
Q4446 aeromonas p
Q4401 aeromonas p
Q4401 aeromonas e
Q95x10 lilium long
Q95x10 pisum sativ
P8236 pisum sativ
P82085 limnodynast
Q15894 homo sapien
Q49493 eopsaltria
Q90493 eopsaltria
Q90493 eopsaltria
Q90493 eopsaltria
Q90493 eopsaltria
Q43960 azorobacter
Q15891 homo sapien
Q83c6 glycine max
Q95c0 hordeum vul
P82222 bombyx mori
Q83c8 hordeum vul
Q82c8 hordeum vul
Q82c8 hordeum vul
Q90c97 homo sapien
Q9bdd9 porilla gor
Q9bdd9 pan proglod
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Best Local :
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071066;
01-AUG-1998
01-AUG-1998
01-DEC-2001
Fusion prote
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092766;
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Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-DEC-2001
Thymidylate
TS.
             Canine
                                                                                                                                                                                                                                                                                                                    proximal coding part of the F gene of wild-type and vacamorbilliviruses.";
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases
EMBL; AF026237; AAC09167.1; -.
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Viruses;
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Liermann H., Harder T.,
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distemper virus.
s; ssRNA negative-strand
                                      -1998 (TrEMBLrel. 07,
-2001 (TrEMBLrel. 19,
protein (Fragment).
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ba; Chordata;
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Pred. No. 1e+0
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P82081;
01-MAY-2000
01-MAY-2000
01-MAY-2000
DYNASTIN 3.
Obermaier
Submitted
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             SEQUENCE FROM N.A.
Obermaier B., Piravandi
Submitted (MAY-1996) to
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NCBI_TaxID=11232
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              the
              Rinke M.;
EMBL/GenBank/DDBJ
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O13591 PRELIMINARY; 2...,
O13591; O1-JAN-1998 (TrEMBLrel. 05, Created)
O1-NOV-1999 (TrEMBLrel. 12, Last sequence update)
O1-OCT-2002 (TrEMBLrel. 22, Last annotation update)
ORF YML337W (Fragment).
Saccharomyces cerevisiae (Baker's yeast).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Sacch
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
STRAIN-Dog #10757/96;
Liermann H., Harder T., Haas L.;
Liermann H., Harder T., Haas L.;
"Genetic analysis of the central untranslated genome proximal coding part of the F gene of wild-type and vorbillivruses.";
Submitted (SEP-1997) to the EMBL/GenBank/DD5J databas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE, AND MASS SPECTROMETRY.

TISSUE=TIBIAL GLAND;

Raftery M.J., Bradford A.M., Bowie J.H., Wallace J.C., Tyler M.J.;

Replides from Australian frogs. The structure of the dynastins from 
"Peplides from Australian frogs. The structure of the dynastins from 
the banjo frogs Limnodynastes interioras, Limnodynastes dumerilii and 
the banjo frogs Limnodynastes interioras, Limnodynastes dumerilii and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Limnodynastes terraereginae (Northen banjo frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Myobatrachi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chem. 46:833-842(1993).
SPECTROMETRY: MW=1236; METHOD=FAB
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100.0%; Pr
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01-NOV-1996 (
01-VOV-2003 (
Aml1 protein
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SEQUENCE
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Submitted
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MEDILINE=96226397, PubMed=8634147;

Levanon D., Bernstein Y., Negreanu V., Ghozi M.C., Bar-Am I.,

Levanon D., Bernstein Y., Lotem J., Groner Y.;

Aloya R., Goldenberg D., Lotem J., Groner Y.;

"A large variety of alternatively spliced and differentially express

mRNAs are encoded by the human acute myeloid leukemia gene AML1.";

DNA Cell Biol. 15:175-185(1996).

EMBL; X90978; CAA62465.1; -.
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[1]
Pfeil B.E., Brubaker C.L., Craven "Phylogeny of Hibiscus and the tri chloroplast DNA sequences of ndhF Syst. Bot. 27:333-350(2002).
EMBL; AF384567; AAM50405.1; -. GO; GO:0009507; C:chloroplast; IEA
                                                                                                                                                                                Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eneurosids II; Malvales; Malvaceae; Malvoideae; Anoda.
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Mammalia; Eutheria;
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                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                               Chloroplast
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                                                                                                                                                              NCBI_TaxID=183227;
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                                                                              Craven L.A., Crisp M.D.;
the tribe Hibisceae (Mal
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SEQUENCE 1
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Q23876;
Q1-NOV-1996
Q1-NOV-1998
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BMBL, K03100; AAA42388.1; -...
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Viruses; Retroid viruses; Retroviridae; Alpharetrovirus.
NCBI_TaxID=11958;
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Eukaryota; Mycetozoa; Dic
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Mycetozoa; Dictyosteliida;
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100.0%; F1
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Pred. No.
4; Mismatc
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Last annotation updat
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Pred. No.
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Pred. No.
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                                                                                           Se+04;
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Best Local S
Matches 3
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Matches 3
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Q35374;
01-NOV-1996
01-NOV-1996
01-UUN-2003
P1 protein (1
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Q28742;
Q1-NOV-1996
Q1-NOV-1996
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STRAIN-stock 172;
MEDLINE=87055241; PubMed=3023187;
MEDLINE=8705241; PubMed=3023187; PubMed=3
                                                                                                                                                                                                                                                                 PIR; 1468
NON TER
SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Alpha-myosin heavy chain (Fragment).
Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NON
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Gene 44:243-253(1986)
                                                                                                                                                                                                                                                                                                                                              ventricular myosin heavy proc. Natl. Acad. Sci. U.; EMBL; K01698; AAA31415.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
MEDLINE=84221901; PubMed=6328491;
Friedman D.J., Umeda P.K., Sinha
Rabinowitz M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUN-2003
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=5888;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Mitochondrion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Paramecium tetraurelia.
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L; M15280; AAA79267.1; -.
GO:0005739; C:mitochondrion;
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                                                                                                       QFMHN 10
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(TrEMBLrel. 01, Last sequence update)
(TrEMBLrel. 24, Last annotation update)
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1266 MW;
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                                                                                                                                                                                                                                                                       MW;
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U.S.A. 81:3044-3048(1984)
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01,
24,
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Last annotation update)
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3; Mismatches
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                                                                                                                                                                                    Score 16; DB 6;
Pred. No. 1e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1D84259D16D046D4 CRC64;
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.1.5e+04;
.0; Indels
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                                                                                                                                                                                                              Length 7;
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SQ SEQUE
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Q40659
ID Q4065
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Q05403
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Matches
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Best Local S
Matches 2
Q40659;
Q40659;
01-NOV-1996
01-NOV-1996
01-OCT-2002
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Q05403;
Q1-NOV-1996
Q1-NOV-1996
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NON TER
SEQUENCE
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Gallus gallus (Chicken).
Gukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Macanathae; Galliformes; Phasianidae; Phasiani
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ol-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 22, Last sequence update)
01-NOV-1996 (TrEMBLrel. 22, Last annotation update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
DNA for ORF'S from chromosome XV (Fragment).
Saccharomyces cerevisiae (Baker's yeast).
Eukaryota, Fungi, Ascomycota, Saccharomycotina, Saccharomycetes,
Saccharomycetales, Saccharomycetaceae; Saccharomyces.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Q8JJ20 PRELIMINARY; PRT; 7 AA. Q8JJ20; CTCATCOT.2002 (TrEMBLrel. 22, Created) 01-OCT-2002 (TrEMBLrel. 22, Last sequence up 01-OCT-2002 (TrEMBLrel. 22, Last annotation Extracellular fatty acid binding protein (Fr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Zumstein E., Pearson B.M., Kalogeropoulos A., "A 29.425 kb segment on the left arm of yeast more than twice as many unknown as known open Yeast 11:975-986(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A. STRAIN=FY1679;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; AF487519; AAL96665.1; -.
NON TER 1 1
SEQUENCE 7 AA; 780 MW; 72CI
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"Cloning and sequencing of 3'
"Submitted (FEB-2002) to the EL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=4932;
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1: geneseqp1980s:*
2: geneseqp1990s:*
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1 SVSEIQFMHNLG 12
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Compugen Ltd.
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Result No.	Score	Query Match	Length	DB	ID	Description
1	50	82.0	11	σį	ABG72607	Abg72607 Parathyro
N	50		12	σν	9	g72608 P
ω	46	'n	11	w	AAY96968	8968 E
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14	36	φ.	9	ω	AAY96966	Para
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16	ω G	4.	10	4.	AAB96932	Aab96932 Rat parat
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invention a	1; Pag	parathyroi isolation ermining bi	003-06	lnik RJ		-2000	-2000	-2002	11087	iffer	apien sp. urus. rofa. famil	para Y hyp	hyroid	3-2003	907;	07 st		31 31 31 31 31 31 31 31 31 31 31 31 31 3
on relate ind isolat	ge 5; 11 <u>r</u>	roid hormone on of antibo bioactive P	66685/06.	, Lavi	ZAHRADNIK R LAVIGNE J R.	; 2000US-	; 2000US-	•	1-A1.	Locat ence 6 /labe	iaris.	chyroid erparat	hormone	(first		andard; [0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
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new antigenic peptide for induci antibodies having an affinity to	ish.	TH) antigenic peptide inducing the shaving an affinity to it, usefulevels in serum, plasma and/or ce					4.			alifiers , Phe		<pre>antigen; hypercalcaemia; osteop ; mouse; rat; bovine; porcine; c</pre>	ic peptide 2-12.			11 AA.	ALIGNMENTS	ABP71484 ABP71485 ABP71483 ABP71483 ABP71483 ABP71483 ABP71482 ABP71482 ABP71482 ABP8547 AAB01864 AAB06891 AAB00891
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Best Local S
Matches 10
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Sus scrofa.
Canis famil:
                                                                              05-DEC-2000; 2000US-00730174
                                                                                                               05-DEC-2000; 2000US-00730174
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; parathyroid hormone; antigen; hypercalcaemia; osteoporosis; primary hyperparathyroidism; mouse; rat; bovine; porcine; canine.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      formed from the N-terminus of parathyroid hormone are; (1) a method for producing antibodies useful PTH levels in a biological sample comprising: (a)
                                                                                                                                                                                 US2002110871-A:
                                                                                                                                                                                                                                                                                                                                                                 Rattus sp.
                            (ZAHR/) ZAHRADNIK R J.
(LAVI/) LAVIGNE J R.
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                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                 /label= Leu, Phe
                                                                                                                                                                                                                                                  /label= Ser, Ala
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Sequence

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The invention relates to a new antigenic peptide for inducing the CC formation and isolation of antibodies having an affinity to it, being CC formed from the N-terminus of parathyroid hormone (PTH). Also included CC are, (1) a method for producing antibodies useful in the determination of CPTH levels in a biological sample comprising: (a) providing at least one CC first peptide antigen comprising appelide fragment of PTH; (b) cc administering the first peptide antigen to a host animal to induce campisher and the interpretation of comprising the first peptide antigen to a host animal to induce comprising antisers produced in the host animal to induce comparisher a produced in the host animal, and (e) selecting compared to produced peptide antigen; (2) an antibody (ab) produced by the method, and (3) test kits and analytical procedures used for the comparisher of the present invention are useful for determining to determining the antigens; antibodies and methods of the present invention, as compared to prior art, have the particular, advantages of possessing greater affinity for PTH, and in particular, are designed to have a novel compared to prior art, have the particular, are designed to have a novel crecognition for anino acid residues extending beyond the first N-terminal PTH residue, and further have negligible cross-reactivity with the large compared to produced in the present sequence represents and primary compared to produced primary compared to produced primary compared to produced primary compared to produced residues are an important parameter in compared to produced produced primary compared to produced primary compared to produced primary compared p
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RESULT 3
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Matches 10
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                                                                                                                                                          Parathyroid hormone N-terminal signaling domain (residues 1-11).
                                                                                                                                                                                               AAY96968;
                                                                                                                                                                                                                AAY96968 standard; peptide;
(GARD/)
(KRON/)
(POTT/)
                                     31-DEC-1998;
                                                       30-DEC-1999;
                                                                                           WO200039278-A2
                                                                                                             Homo sapiens
                                                                                                                              parathyroid hormone; N-terminal; signaling domain; bone mass; reformation; resorption; remodeling; tetherl; osteoporosis.
) GARDELLA T J.
) KRONENBERG H M.
) POTTS J T.
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90.9%;
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Zahradnik RJ,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        regular injections
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                                                                                                                                                                                                                                                                                                                    diagnosis;
                                                                                                                                                                                                                                                                                                                                                              Human
                                                                                                                                                                                                        28-SEP-1994;
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                                                     antigenic peptide(s) frerated using them, able
                                                                                             1996-180391/19
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                                                                                                                                                                                                                                                                                                                                                            parathyroid
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                                                                                                                                                                                                                                                                                                                                   hormone; hPTH;
                                                                                                                                                                                                                                                                                                                                                            hormone antigenic peptide hPTH 1-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                              peptide;
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81.8%;
                                                      from
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Potts JT,
                                                     human parathyroid hormone - and antibodies distinguish between active and inactive
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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                                                                                                                                                                                                                                                                                                                                     antigen; alpha-helix; immunogen;
                                                                                                                         Maegerlein
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Page 4;

German

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Matches

Similarity

73.8%;

Score 45; Pred. No.

Length 10;

Mismatches 0 DB 3; .082;

Indels

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Conservative

SVSEIQEMHN 10 SVSEIQLMHN

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Sequence

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Best Local :
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                                    The present sequence represents a parathyroid hormone (pTH) fragment, comprising amino acids 1-10. Parathyroid hormone fragments of the invention have lipolysis stimulating activity (especially when topically administered). The lipolytic activity of the peptides is enhanced when they are chemically modified to increase their lipophilicity. The peptides are used in cosmetic or dermatological compositions for skin care. They are especially used for slimming treatment of excessive weight in the thighs and hips, in the treatment of cellulite and for skin firetree.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Parathyroid hormone; lipolysis; cosmetic; dermatological; skin care; slimming treatment; cellulite; skin firming.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY68767;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY68767 standard;
                                                                                                                                                                                                      in topically applied cosmetic composexcessive weight in hips and thighs.
                                                                                                                                                                                                                     New parathyroid hormone fragment peptides, used as lipolysis stimulants in topically applied cosmetic compositions for slimming treatment of
                                                                                                                                                                                                                                                                                                                                                                                07-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                             27-JAN-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Amino acids 1-10 of a parathyroid hormone (pTH)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-MAY-2000
                                                                                                                                                                         Claim 1; Page 8; 18pp; French.
                                                                                                                                                                                                                                                                WPI; 2000-171243/15.
                                                                                                                                                                                                                                                                                            Lintner
                                                                                                                                                                                                                                                                                                                       (SEDE-) SEDERMA
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90.0%;
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Pred. No. 0.082;
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RESULT 6
AAB86219
ID AAB8
RESULT 7
ABR44166
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                                                                                                                                                                                                                                                                                                                                                                                                             CC This invention describes a novel method for determining (M1) the content CC of active parathyroid hormone (A) by treating a sample with (i) antibody CC (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), CC and including the N-terminal residue and (ii) antibody (Ab2) that CC recognizes an epitope within the receptor-binding site of (A). The number CC of molecules that react with both antibodies is determined and used to CC calculate the content of physiologically active (A). The method is used CC of diagnosis of calcium-metabolism disorders, osteopathy, or hypo- or CC phyper-parathyroidism. The method (unlike known assays) recognizes that CC (i) some fragments of (A) shorter than the complete (84 aa) peptide are CC cactive and (ii) that apparently intact peptide may be biologically CC are antagonistic (these have the receptor-binding site but lack the N-CC terminus). It thus provides a true measure of the content of CC physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This CC sequence represents a peptide fragment used to illustrate the method of the content of the content of the content of the content of content of the content of content 
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Best Local S
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibodies reactive with different epitopes.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-SEP-2001
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                   ABR44166
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                                                             ABR44166 standard; peptide; 10
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                                                                                                                                                                                                                                                                   Similarity
9; Conserv
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                                                                                                                                                                             SVSEIQLMHN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunogenic peptide
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Pred. No. 0.08
0; Mismatches
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                                                                                                                                                                                                                                                                                        DB 4;
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RESULT 8
AAB86225
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to a fusion peptide (Tat-hPTHDP), where self cell penetrating Tat peptide is bound to human parathyroid hormone-derived peptide (hPTHDP). The fusion peptide is useful as a component of skin slimming cosmetic composition. The fusion peptide does not cause irritation, easily and safely penetrates into integument and endotheliu does not cause skin disease and has superior lipolysis effects, and is durable. The present sequence represents a human parathyroid hormone (hPTH) fragment that can be used to construct the fusion peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel fusion peptide comprising self cell-penetrating Tat peptide to human parathyroid hormone-derived peptide, useful as component slimming cosmetic composition.
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lipolysis; human; hPTH.
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                                                                                                                                                                                                                                 Parathyroid hormone; hPTH; immunogenic peptide; human; epitope diagnosis; calcium-metabolism disorder; osteopathy; antagonist hypo-parathyroidism; hyper-parathyroidism.
                                                                                                                                                                                                                                                                                                                                          Human parathyroid hormone immunogenic peptide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 5; Page 6; 32pp; English.
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15-MAR-2002; 2002KR-00014062
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB86225
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9; Conserv
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Pred. No. 0.082;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 10;
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17-DEC-1999;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            of active parathyroid hormone (A) by treating a sample with (i) antibody (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), and including the N-terminal residue and (ii) antibody (Ab2) that recognizes an epitope within the receptor-binding site of (A). The number of molecules that react with both antibodies is determined and used to calculate the content of physiologically active (A). The method is used for diagnosis of calculamemetabolism disorders, osteopathy, or hypo- or hyper-parathyroidism. The method (unlike known assays) recognizes that (i) some fragments of (A) shorter than the complete (84 aa) peptide are active and (ii) that apparently intact peptide may be biologically cinactive, and also takes into account the fact that some fragments of (A) are antagonistic (these have the receptor-binding site but lack the N-terminus). It thus provides a true measure of the content of physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This sequence represents a peptide fragment used to illustrate the method of
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Best Local S
Matches 8
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                                                                                                                                                                                                                                              Human parathyroid diagnosis; active
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 9
                                                                                                                                                             04-APR-1996.
                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 3; 10pp; German
                WPI; 1996-180391/19
                                            Adermann K,
                                                                                                    28-SEP-1994;
                                                                                                                                 28-SEP-1994;
                                                                                                                                                                                          DE4434551-A1
                                                                                                                                                                                                                                                                                           Human parathyroid hormone antigenic peptide hPTH 1-9.
                                                                                                                                                                                                                                                                                                                        06-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                                AAR91645 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    invention describes a novel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
8; Conserv
                                                                         FORSSMANN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VSEIOFMEN 10
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                                            Forssmann
                                                                                                                                                                                                                                                                                                                       (first entry)
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                                                                                                     94DE-04434551
                                                                                                                                 94DE-04434551
                                                                                                                                                                                                                                                hormone; hPTH; antigen; alpha-helix; hPTH 1-37.
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                                                                                                                                                                                                                                                                                                                                                                                peptide;
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Pred. No. 1.4e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     method for determining (M1) the content by treating a sample with (i) antibody
                                            Maegerlein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .4e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 9;
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                                                                                                                                                                                                                                                                 immunogen,
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RESULT 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence is a specific example of claimed immunogenic peptides having a sequence from hPTH(1-37) which includes the N-c terminal alpha-helical region and/or the non-structured region of hormone. Antibodies and their binding fragments generated by injection and with the peptides are useful as diagnostic reagents for determination of biologically active hPTH(1-37)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (GARD/)
(KRON/)
(POTT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gardella TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-DEC-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Parathyroid hormone N-terminal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 2; Page 4;
                                                                                                                                                                                                                                                                                                                                                                                                                            2000-452384/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             parathyroid reformation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ) KRONENBERG H M.
) POTTS J T.
JIPEDDOTTS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JUEPPNER
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kronenberg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hormone; N-t; resorption;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-terminal; signaling ion; remodeling; tether
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 39; DB
Pred. No. 1.4e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Jueppner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             naling domain;
tether1; oste
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Compounds of the structure or formula S-(L)_n-B, R_1-S-(L)_n-R or S-(L)_R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH), L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4); B is a C-terminal binding portion of pTH(1-34) or a pTH-related protein (PTH-TP) (1-34); R_1 is tip PTH-1 receptor signal sequence; and R is (a portion of) PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAu in a mammalian cell having PTH-1 receptors, or screening for a peptide of the sequence of th

binding R_1 is the

(where

increasing cAMP or a peptide or

New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.

Claim 11;

Page 93; 119pp;

English.

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Query Match
                    Sequence 9
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63 . 98

Score

39;

DB 4.

Length

This invention describes a novel method for the preparation of peptides CC (II) that contain both cyclic and linear peptide fragments comprises CC sequential reaction of a resin-bound linear fragment with the cyclic fragment in N-protected form and optionally other linear fragments. The CC bind to hPTH receptors and act as agonists or antagonists of hPTH. The CC method is particularly used to prepare cyclic peptide analogs of CC parathyroid hormone (PTH) or PTH-related peptides which are useful for CC treating diseases that respond to treatment with agents that bind to PTH CC receptors (with or without activation of adenylyl cyclase activity), e.g. CC hyper- or hypo-calcemia, osteoporosis, osteopenia, hyper- or hypo-calcemia, osteopenia, hyper- or hypo-c

The (II)

Disclosure; Page 75;

85pp; English.

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RESULT 11
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                                         This invention describes a novel method for determining (M1) the content CC of active parathyroid hormone (A) by treating a sample with (i) antibody CR (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), CC and including the N-terminal residue and (ii) antibody (Ab2) that CR recognizes an epitope within the receptor-binding site of (A). The number of modecules that react with both antibodies is determined and used to calculate the content of physiologically active (A). The method is used for diagnosis of calcium-metabolism disorders, osteopathy, or hypo- or typer-parathyroidism. The method (unlike known assays) recognizes that (i) some fragments of (A) shorter than the complete (84 aa) peptide are active and (ii) that apparently intact peptide may be biologically cinactive, and also takes into account the fact that some fragments of (A) are antagonistic (these have the receptor-binding site but lack the N-cerminus). It thus provides a true measure of the content of physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This sequence represents a peptide fragment used to illustrate the method of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   non-peptide PTH (claimed). The new compound can be administered by inhalation unlike the large native PTH or PTHrP which avoids the native properties to treat osteoporosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist;
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                                                                                                                                                                                                                                                                                                                                                                                 Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antiboc reactive with different epitopes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Armbruster FP
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8; Conservative
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                                                                                                                                                                                                                                                                                                                                               10pp;
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RESULT 12
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Matches 8; Conserval
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Convergent synthesis of peptides for treating e.g. bone disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sledeski AW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15-APR-1998;
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The invention relates to a novel parathyroid hormone (PTH) peptide (CAB01859) and parathyroid hormone-related peptide (PTHrP, AAB01860), and biologically active derivatives thereof (AB01857-B01858, AB01861). The peptides of the invention are at least 85% identical to the B01869). The peptides of the invention are at least 85% identical to the generic peptide of the formula: X1-Val-Ser-Glu-X2-Gln-Leu-X3-H18-X4-X5-CGly-Lys-X6 (AAB01857) where: X1 is Ser or Ala: X2 is 11e or Lys: X3 is Ser or Ala: X2 is 11e or Lys: X6 is His or Ser: Met, Leu or Nle: X4 is Asn or Asp; X5 is Leu or Lys: X6 is His or Ser: Met, Leu or Nle: X4 is Asn or Asp; X5 is Leu or Lys: X6 is His or Ser: Met, Leu or Nle: X4 is Asn or Asp; X5 is Leu or Lys: X6 is His or Ser: Met, Leu or Nle: X4 is Asn or Asp; X5 is Leu or Lys: X6 is His or Ser: Met, Leu or Nle: X4 is Asn or Asp; X5 is Leu or Lys: X6 is His or Ser: Met, Leu or Nle: X4 is Asn or Asp; X5 is Leu or Lys: X6 is His or Ser: Met, Leu or Nle: X4 is Asn or Asp; X5 is Leu or Lys: X6 is His or Ser: Met, Leu or Nle: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Ser: Met, Leu or Lys: X6 is His or Leu or Lys: X6 is His or Leu or Lys: X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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that encode them, useful for
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) KRONENBERG H M.
) POTTS J T.
) JUEPPNER H.
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Pred. No. 1
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) KRONENBERG H M.
) POTTS J T.
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1; Mismatches
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1.4e+06;
ches 1;
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compounds of the structure or formula S-(L)_n-B, R_1-S-(L)_n-R or S-(L)_R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (when n is a positive integer 1-10 (preferably 4)); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-rP) (1-34); R 1 is t PTH-1 receptor signal sequence; and R is (a portion of) PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases

binding R 1 is the

(where

s-(L)_n

New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.

WPI; 2000-452384/39

Claim 4; Page 92; 119pp; English

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RESULT 15
AAB01863
       The invention relates to a novel parathyroid hormone (PTH) peptide (AAB01859) and parathyroid hormone-related peptide (PTH-P; AAB01860), and biologically active derivatives thereof (AAB01857-B01859, AAB01861-CC B01859). The peptides of the invention are at least 85% identical to the generic peptide of the formula: X1-Val-Ser-Glu-X2-Gln-Leu-X3-His-X4-X5-CC Gly-Lys-X6 (AAB01857) where: X1 is Ser or Ala; X2 is Ile or Lys; X3 is CC Met, Leu or Nle; X4 is Asn or Asp; X5 is Leu or Lys; X6 is His or Ser; CC provided that the peptide is not PTH-P(1-14). The peptides of the consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C- terminal derivatives. PTH is a major regulator of calcium homeostasis, CC and is necessary for the normal function of the gastrointestinal, reurological system, neuromuscular and cardiovascular systems. CC and to the recently identified PTH-2 receptor. PTH has a potent anabolic effect on the skeleton, and mediates calcium reabsorption, enhances CC phosphate clearance and vitamin D synthesis in the kidney. A homologous
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Parathyroid hormone peptide; PTH; PTH-related peptide; PTHrP; calcium homeostasis; PTH-1 receptor; PTH-2; vitamin D synthesis; bone synthesis; agonist; osteoporosis; non-parenteral delivery.
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                                                                                                                                                                                                                                                                                                                       Disclosure; Page 26; 73pp; English
                                                                                                                                                                                                                                                                                                                                                      Parathyroid hormone (PTH) peptides, acids that encode them, useful for t
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Pred. No. 1.4e+06;
1; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                          treating
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calls of protein hormones, the PTH-related proteins (PTHrP) mimic some of characteristics and skeletal actions of PTH, and also bind to the PTH-1 creeptor. They do not bind to the PTH-2 receptor. The peptides of the invention are either agonists of PTH-1 and PTH-2 receptors (AAB01858, AAB01861-B01866) or are PTH-1/PTH-2 receptor antagonists. (AAB01858, B01869) PTH-1/PTH-2 receptor agonists are useful for the treatment of conditions characterised by a decrease in bone mass, such as conceoporosis. PTH-1/PTH-2 receptor antagonists are useful for treating medical disorders that arise from excessive or altered action of the PTH-1/PTH-2 receptor. Detectably labelled peptides of the invention are also useful in the determination of rates of bone formation, bone resorption and/or bone remodelling in a patient. The peptides of the invention are "minimised" versions of PTH or PTH-P which are inexpensive to prepare by conventional synthetic chemistry, and can be delivered to a prepare by C non-parenteral routes. Sequences AAB01858 and AAB01866 represent convention are receptor agonists.
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Sequence 9 AA;

Matches Query Match Best Local 6 Similarity Conservative 54.18; Score 33; DB 3; Pred. No. 1.4e+06; 2; Mismatches 1 DB 3; Length 9; Indels 0, Gaps 0

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Search completed: May 18, 2004, 10:01:54 Job time : 46.75 secs

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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

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Copyright (c) 1993 - 2004 Compugen Ltd.
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equence 41, Ap	equence 12,	quence 4, Appl	equence 4, Appl	equence 2058, 7	equence 20	equence 14	Sequence 1385, Ap	equence 12	equence 10	equence 95	equence 40	equence 33	equence 12	equence 26	equence 50	equence 16	equence 14	equence 8,	equence 78,	equence 73,	equence 6,	quence 55,	equence 45,	equence 2,	equence 52	equence 11	equence 8,	ence 4,	100

ALIGNMENTS

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OTHER INFORMATION: Induces Formation of Antibodies and Isolates US-09-730-174A-4
Sequence 6, Application US/09730174A
Patent No. US20020110871A1
GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid FILE REFERENCE: IMUNB-001A
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Patent No. US2002011871A1

GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.

TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having ITILE OF INVENTION: Selective Binding Specificity to Bicactive Intact Parathyroid H

FILE REFERENCE: INUNE-001A

CURRENT FILING DATE: 2000-12-05

NUMBER OF SEQ ID NO 4

LENGTH: 12

TYPE: PET

ORGANISM: Artificial Sequence
FEATURE:
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US-09-730-174A-3
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CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 6
LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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SEQ ID NO 2
LENGTH: 11
TYPE: PRT
ORGANISM: Artificial Sequence
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Patent No. US20020110871A1
GENERAL INFORMATION:
                                                                                                                            CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 3
LENGTH: 12
TYPE: PRT.
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                  Sequence 3, Application US/09730174A Patent No. US20020110871A1 GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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Best Local :
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Best Local
               Best Local Similarity 91. Matches 11; Conservative
                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                             TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies
TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parath
FILE REFERENCE: IMUNE-001A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: IMUNE-001A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having
TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid;
                                                                                   -09-730-174A-3
                                                                                                 OTHER INFORMATION:
                                                                                                                   FEATURE:
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11; Conserv
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11; Conserv
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llarity 100.0%;
Conservative (
                                                                                                 Induces Formation of Antibodies and Isolates
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91.7%;
                               90.2%;
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Pred. No. 0.00
1; Mismatches
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              Score 55; DB 9; Pred. No. 0.0017; 0; Mismatches
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                                                 Length 12
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US-09-730-174A-5
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Best Local S
Matches 10
                                               Sequence 1, Application US/10168185
Publication No. US20030175802A1
GENERAL INFORMATION:
APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local
   APPLICANT: Schmidt-
APPLICANT: Roth, He
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                    l Similarity 90.
10; Conservative
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Missbichler, Albert
Schmidt-Gayk, Heinrich
Roth, Heinz-Jurgen
NVENTION: Method for Determining Parathormone
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Sequence 5, Application US/09730174A

Patent No. US20020110871A1

GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.

TITLE OF INVENTION: Antibodies and Peptide Antigens for Pro-
TITLE OF INVENTION: Selective Binding Specificity to Bioac
FILE REFERENCE: IMUNE-001A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
                                                                                                                                                 APPLICANT: Zahradnik, R.J.

ITTLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having ITTLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid H FILE REFERENCE: IMUNE-001A

CURRENT APPLICATION NUMBER: US/09/730,174A

CURRENT FILING DATE: 2000-12-05

NUMBER OF SEQ ID NOS: 12

SEQ ID NO 1

LENGTH: 11

TYPE: PRT
                                                                           , OTHER INFORMATION: Induces Formation of Antibodies and Isolates US-09-730-174\mbox{A-1}
                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1, Application US/09730174A Patent No. US20020110871A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                              ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Induces Formation of Antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             85.2%;
                 90.9%;
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Pred. No. 0.00
1; Mismatches
0
               Score 51;
Pred. No.
  Mismatches
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                 DB 9;
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                                  Length 11
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  Gaps
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APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Method for Determining Parathormone
FILE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: MUMBER: US/10/168,185
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: COT/EP00/12911
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION DATE: 1999-12-17
NUMBER OF SEG ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NOS: 1
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CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
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Best Local Similarity
Thes 9; Conserve
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US-10-168-185-7
             Sequence 2, Application US/10168185
Publication No. US20030175802A1
GENERAL INFORMATION:
APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 7, Application US/10168185
Publication No. US20030175802A1
GENERAL INFORMATION:
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TYPE: PRT
ORGANISM: Homo sapiens
CURRENT APPLICATION NUMBER: US/10/168,185
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TYPE: PRT
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88.9%;
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Pred. No. le+06;
0; Mismatches
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Pred. No. 0.098;
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APPLICANT: Ambruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION WIMBER: PGT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
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US-10-192-673-6
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                                                                                                                                                                                                                                               Sequence 6, Application US/10192673; Publication No. US20030166838A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 8, Application US/10168185 Publication No. US20030175802A1 GENERAL INFORMATION:
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Best Local :
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PRIOR APPLICATION NUMBER: PCT/BE00/12911
PRIOR PILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEO ID NOS: 11
SOPTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Hormone-Related Peptide (PTHP) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHP)
TILE REFERENCE: 0609-4570002
CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: US/10/192,673
CURRENT FILING DATE: 2002-07-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 8
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
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Pred. No. 1e+06;
0; Mismatches
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Pred. No.
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APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Ports, John T.
APPLICANT: Ports, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR APPLICATION NUMBER: US. 60/105,530
PRIOR FILING DATE: 1998-10-22
PRIOR FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
NUMBER OF SEQ ID NOS: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-10-192-673-6
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US-10-192-673-7
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                                                                                                                                                                                                                                         RESULT 13
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Publication No. US20030166838A1
GENERAL INFORMATION:
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SEQ ID NO 6
                                                                                                GENERAL INFORMATION:
APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
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Best Local
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Publication No. US20030175802A1
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Best Local (
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APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE: OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence
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7; Conservative
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77.8%;
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Pred. No. 1e+06;
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Pred. No. le+06;
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APPLICANT: Juppner, Harald
ITILE OF INVENTION: Bloactive Peptides and Peptide Derivatives of
ITILE OF INVENTION: Bloactive Peptides and Peptide Derivatives of
ITILE OF INVENTION: Hormone-Related Peptide (PTHP)
ITILE OF INVENTION: Hormone-Related Peptide (PTHPP)
ITILE REFERENCE: 0609,4570002
CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR FILING DATE: 1999-10-20
PRIOR FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PSI/PPO0/12911
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
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Best Local S
Matches 6
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Best Local S
Matches 7
                                                                                                 Sequence 61, Application US/10033741
Publication No. US20030049640A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 10, Application US/10192673 Publication No. US20030166838A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
APPLICANT: Herman, et al.
TITLE OF INVENTION: Proteins, Genes and T
TITLE OF INVENTION: Response
FILE REFERENCE: 9195-079
CURRENT APPLICATION NUMBER: US/10/033,741
                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
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6; Conserv
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87.5%;
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                                                              Genes and Their Use For Diagnosis and Treatment
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Pred. No. 1e+06;
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Pred. No. 1e+06;
1; Mismatches
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Scoring table:
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1: /cgn2 6/ptodata/2

2: /cgn2 6/ptodata/2

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4: /cgn2 6/ptodata/2

5: /cgn2 6/ptodata/2

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Query
Match
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61
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                                                                               Copyright
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/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/BCTUS_COMB.pep:*
/cgn2_6/ptodata/2/iaa/pcTUS_COMB.pep:*
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US-09-442-989-31
US-08-817-547A-1
US-08-817-547A-2
US-09-421-379-6
US-09-421-379-7
US-08-817-547A-2
US-08-817-547A-4
US-08-817-547A-4
US-08-817-547A-4
US-08-817-547A-4
US-08-1482-528-42
US-08-177-109A-5
US-08-177-109A-1
US-08-17-547A-1
US-08-817-547A-1
US-08-817-547A-1
US-08-817-547A-1
US-08-817-547A-1
US-08-817-547A-1
US-08-817-547A-1
US-08-817-547A-7
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Sequence 2, Appli
Sequence 2, Appli
Sequence 6, Appli
Sequence 7, Appli
Sequence 10, Appli
Sequence 42, Appl
Sequence 42, Appli
Sequence 5, Appli
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Sequence 56,
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US-09-442-989-31
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US-08-428-257A-14	US-08-555-579-2	- 18	1	US-09-064-750-1	US-09-556-605-61	-09-	US-09-434-122-49	US-09-393-385B-108	US-09-344-050-12	US-09-280-028-12	US-08-836-561-49	53A-	US-08-483-528B-97	US-08-438-562-9	US-08-116-778E-9	PCT-US95-06613-26	'n
Sequence 14, Appl	-	3, App	,	equence 1, App1	equence 61,	e 3, Appı	e 49, App	e 108, Ap	e 12,	e 12, App	equence 49;	e 12,	equence 97,	equence 9, F	equence	equence 26,	equence 26, App

ALIGNMENTS

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; FEATURE:
, NAME/KEY: MOD RES
; LOCATION: (8)
; OTHER INFORMATION: N1e
US-09-442-989-31
                                                                                                                                            PEATURE:
NAME/KEY: PEPTIDE
LOCATION: (4)
OTHER INFORMATION: (
FEATURE:
                                                                                                                                                                                       NAME/KEY: PEPTIDE
LOCATION: (3)
OTHER INFORMATION: Ser(OtBu)
                                             OTHER INFORMATION:
                                                     NAME/KEY: PEPTIDE (10)
                                                                                LOCATION: (9)
OTHER INFORMATION: His(Trt)
                                                                                                                  OTHER INFORMATION:
                                                                                                                                                                                                                            OTHER INFORMATION: FMOC-Ala
                                                                                                                                                                                                                                    NAME/KEY: PEPTIDE LOCATION: (1)
                                                                                                  NAME/KEY: PEPTIDE
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                                                                                                                  Gln (Trt
                                                                                                                                                      Glu (OtBu)
                                             Asn (Trt)
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RESULT 2 US-08-817-547A-1

GENERAL INFORMATION:
APPLICANT: Adermann, Knu
APPLICANT: Hock, Dieter

Adermann, Knut

STREET: 191. CITY: Atlanta
STATE: Georgia
STATE: Georgia

CORRESPONDENCE ADDRESS:

ADDRESSEE:

```
ZIP: 30303
ZIP: 30303
ZIP: 30303
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTEN PC-DOS/MS-DOS
SOFTWARE: PATENTEN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 07826-0007
FELEPRENCE/DOCKET NUMBER: 07826-0007
TELEPRENCE/DOCKET NUMBER: 07826-0007
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Best Local Similarity 75.0%;
Matches 9; Conservative
                                                                                                                         Sequence 2, Application US/08817547A Patent No. 6030790
APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: Jones & Askew, LLP
191 Peachtree Street, 37th Floor
                                                                                                                                                                                                                                                                                                                                                                                         73.8%; Score 45; DB 3; Length 10; 90.0%; Pred. No. 0.017; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 46; DB 4; Pred. No. 0.013; 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
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                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Note: John T.
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Hormone-Related Peptide (PTHP)
FILE REFERENCE: 0609.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: synthetic
; OTHER INFORMATION: peptide
US-09-421-379-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: po

HYPOTHETICAL: no

ANTI-SENSE: no

US-08-817-547A-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 8; Conserv
                                                                                                                                                 SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 6
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 6, Application US/09421379 Patent No. 6495662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE:

PRIOR APPLICATION DATA:

PRIOR APPLICATION UNWER: PCT/EP95/03757

APPLICATION UNWER: PCT/EP95/03757

FILING DATE: 29 SEPT 1994

ATTORNEY/AGENT INFORMATION:

NAME: FROST, ROGER T.

REGISTRATION NUMBER: 22,176

REGISTRATION NUMBER: 07826-0007

TELECOMOUNICATION INFORMATION:

TELEPHONE: 404-818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 30303
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A
                                                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 191 Pe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
STRANDEDNESS: un
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SVSEIQFMH 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     63.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 39; DB 3; Pred. No. 3e+05; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 9;
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US-08-817-547A-2

GENERAL INFORMATION:

g S

Query Match Best Local S Matches 9

Similarity 9; Conserv

Conservative

SVSEIQFMHN 10

TOPOLOGY: unkno MOLECULE TYPE: pe HYPOTHETICAL: no ANTI-SENSE: no

LENGTH: 10 amino acids TYPE: amino acid STRANDEDNESS: unknown

unknown peptide

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Sequence 7, Application US/09421379

Patent No. 6495662

GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Fotts, John T.
APPLICANT: Fotts, John T.
APPLICANT: Wippner, Harald
ITITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
ITITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
ITITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
ITITLE OF INVENTION: Wormone-Related Peptide (PTHP)
FILE REFERENCE: 0609.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: U.S. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0

SEQ ID NO. 7
                                                                                                                                                                                                                                                             Sequence 3, Application US/08817547A
Patent No. 6030790
GENERAL INFORMATION:
APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Se
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
ADDRESSEE: Jones & Askew, LLP
STREET: 191 Peachtree Street, 37th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide
US-09-421-379-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 5
US-09-421-379-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-817-547A-3
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Best Local
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM FO compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A FILING DATE: PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                CITY: Atlanta
STATE: Georgi
                                                                                                                                                                                    ZIP: 30303
                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 AVSEIQLLH 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 SVSEIQFMH 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity 66.7%;
6; Conservation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SVSEIQFMH 9
                                                                                                                                                                                                        Georgia
Y: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 33; DB 4;
Pred. No. 3e+05;
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 36; DB 4;
Pred. No. 3e+05;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                       the hPTH Sequence
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                                                                                                         RESULT 8
US-08-140-137A-42
Sequence 42, Application US/08140137A
Patent No. 5817617
GENERAL INFORMATION:
APPLICANT: TUOMANEN, ELAINE
APPLICANT: MASURE, H. R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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Sequence 10, Application US/09421379

Sequence 10, Application US/09421379

Patent No. 6495662

GENERAL INFORMATION:

APPLICANT: Excatella, Thomas J.

APPLICANT: Kronenberg, Henry

APPLICANT: Kronenberg, Henry

INTILE OF INVENTION: Bloactive Peptides and Peptide Derivatives of TITLE OF INVENTION: Harald

TITLE OF INVENTION: Harathyroid Hormone (PTH) and Parathyroid TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)

FILE REFERENCE: 0609.4570001

FILE REFERENCE: 0609.4570001

CURRENT APPLICATION NUMBER: US/09/421,379

CURRENT APPLICATION NUMBER: US. 60/105,530

EARLIER APPLICATION NUMBER: US. 500/105,530

EARLIER APPLICATION NUMBER: US. 500/105,530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
FYPE: amino acid
STRANDENNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: peptide
HYPOTHETICAL: no
ANTI-SENSE: no
US-08-817-547A-3
                                                                                                                                                                                        ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-09-421-379-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-421-379-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity Matches 7; Conserv
                                                                                                                      Query Match
Best Local Similarity
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FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 07826-0007
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 404-818-3799 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                       1 SVSEIQFMH 9
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                                                                                                 <u>ه</u>
SVSEHQLLH
                                                                                                 Conservative
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                                                                                                                      47.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Mismatches
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Pred. No. 3e+05;
                                                                                                 Score 29; DB 4;
Pred. No. 3e+05;
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 8;
                                                                                                                                              Length 9;
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BRUCE E.; WETTENHALL, RICHARD E.H.

TITLE OF INVENTION: PROTEIN ACTIVE IN HUMORAL
;HYPERCALCEMIA OF MALIGNANCY-PTHRP
NUMBER OF SEQUENCES: 4
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/715,280
FILING DATE: 14-JUN-191
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 199,235
FILING DATE: 09-MAY-1988
APPLICATION NUMBER:
FILING DATE: 09-MAY-1988
APPLICATION NUMBER:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
""" hes 5; Conserva
뮍
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 133521
INFORMATION FOR SEQ ID NO: 42
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acid
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 9
                                                                                                                                                                                                                                                                                                                                                                                                                Patent No. 5460978
, APPLICANT: MARTIN, THOMAS J.; MOSELEY, JANE M.; KEMP,
                                                                                                                                                                                 SEQ ID NO:3:
                                                                                       Query Match
Best Local
                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 600-1-096
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201 343-1684
TELEPAX: 201 343-1684
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/140,137A
FILING DATE: 27-MAY-1994
CLASSIFICATION: 424
CLASSIFICATION: 424
APPLICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: 41. STREET: Hackensack
CITY: Hackensack
CTATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide
DESCRIPTION: CARD peptide segence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Klauber & Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 202
TELEFAX: 133521
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                             LENGTH: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 VSEIQFMH 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 07601
                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ISEEOFVH 12
                                   SVSEIQFMHN 10
AVSEHQLEHN 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   411 Hackensack Avenue
                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         47.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ANALOGS OF ENDOTHELIAL LEUKOCYTE ADHESION MOLECULE (ELAM)
                                                                                       45.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             49
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 29; DB 2; Length 12; Pred. No. 23; 2; Mismatches 1; Indels
                                                                         1; Mismatches
                                                                                       Score 28; DB 6; Length 11;
Pred. No. 32;
                                                                         Indels
                                                                         0
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                                                                         Gaps
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RESULT 10
US-08-817-547A-5
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; HYPOTHETICAL: no
; ANTI-SENSE: no
US-08-817-547A-5
                                                                                                                                                                                                                                                           US-08-817-547A-4
                                                                                                                                                                                                                                                                            RESULT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 5, Application US/08817547A Patent No. 6030790 GENERAL INFORMATION:
                                                                                                                                                                                                 Sequence 4, Application US/08817547A Patent No. 6030790 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/817,547A
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION UNMBER: PCT/EP95/03757
APPLICATION TOWNER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: PROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 07826-0007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
ENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Adermann, .....APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
APPLICANT: Magerlein Merkus
TITLE OF INVENTION: Peptides
7 770TENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Jones & Askew, LLP
STREET: 191 Peachtree Street,
                                                                                                          APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 1...
STREET: 1...
CTTY: Atlanta
CAOTGir
CITY: Atlanta
STATE: Georgia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                   SVSEIQ 6
                                                                                                                                                                                                                                                                                                                                SVSEIQ 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Jones & Askew, LLP
191 Peachtree Street, 37th Floor
                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 404-818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                peptide
                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Pred ...
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                       42.6%; Score 26;
100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07826-0007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         from the hPTH Sequence
                                                                                                                               from the hPTH Sequence
                                                         37th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                                         3e+05;
                                                                                                                                                                                                                                                                                                                                                                                                        0;
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0

COUNTRY: U ZIP: 30303

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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Fotts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: U.S. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 12
US-09-421-379-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-817-547A-4
                                                                                                                                                    US-09-421-379-8
                                                    Query Match
Best Local Similarity
Matches 5; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 8, Application US/09421379 Patent No. 6495662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/
APPLICATION NUMBER: PC94
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 0782:
TELECOMMUNICATION INFORMATION:
TELECHONE: 404.818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                            FEATURE: OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
HYPOTHETICAL: n
ANTI-SENSE: no
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A
FILING DATE:
PRICE ATTOMATE
                                                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 100 nes 6; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: unl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
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                                 SVSEIQFMH 9
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                                                                         Conservative
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100.0%; Fred. No.
                                                                                           42.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  07826-0007
                                                                                       Score 26; DB 4;
Pred. No. 3e+05;
                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 3;
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                                                                         2; Indels
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US-08-482-228-42
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                                                                                                                                                                                                                                                                                           US-08-482-528-42
                                                                                                                                                                                                                                                      Sequence 42, Application US/08482528 Patent No. 6017719
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Tseng-Law, Janet
APPLICANT: Kobori, Joan A.
APPLICANT: Al-Abdaly, Fahad A.
APPLICANT: Guillermo, Roy
                                               APPLICANT: Guillermo, Roy
APPLICANT: Helgerson, Sam L.
APPLICANT: Deans, Robert J.
TITLE OF INVENTION: POSITIVE AND POSITIVE/NEGATIVE CELL
TITLE OF INVENTION: SELECTION MEDIATED BY PEPTIDE RELEASE
NUMBER OF SEQUENCES: 215
CORRESPONDENCE ADDRESS:
ADDRESSEE: Janice Guthrie, Ph.D.
STREET: P.O. Box 15210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                           5 SSVTFMH 11
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MOLECULE TYPE: peptide US-08-482-228-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (714) 553-1952
INFORMATION FOR SEQ ID NO: 42
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No. 5700 Patent Patent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: USA
ZIP: 92713-5210
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,228
FILING DATE: 07-UN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Githy: Tailor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 42, Application US/08482228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: TSeng-Law, Janet
APPLICANT: Kobori, Joan A.
APPLICANT: Kobori, Joan A.
APPLICANT: Al-Abdaly, Fahad A.
APPLICANT: Guillermo, Roy
APPLICANT: Helgerson, Sam L.
APPLICANT: Deans, Robert J.
TITLE OF INVENTION: POSITIVE AND POSITIVE/NEGATIVE CELL
TITLE OF INVENTION: SELECTION MEDIATED BY PEPTIDE RELEASE
NUMBER OF SEQUENCES: 215
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Guthrie, Janice
REGISTRATION NUMBER: 35,170
REFERENCE/DOCKT NUMBER: IT-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (714) 440.5353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Janice Guthrie, Ph.D. STREET: P.O. Box 15210
CITY: Irvine
STATE: California
                                                                                                                                                                  Match 41.0%;
Local Similarity 57.1%;
3 SEIQFMH 9
                                                                                                                                                                                                                                                                                                                                                                                                                                       amino acid
)GY: linear
                                                                                                              Conservative
                                                                                                        Score 25; DB 2;
Pred. No. 1.3e+0
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IT-4630CIP3
                                                                                                  3e+02;
2;
                                                                                                                                                                                                                      Length 12;
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CITY: Irvine STATE: California

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TELEPHONE: (714) 440-5353;
TELEFAX: (714) 553-1952;
INFORMATION FOR SEQ ID NO: 42
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-482-528-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 15
US-08-177-109A-56
(Sequence 56, Application US/08177109A
; Patent No. 5869615
; Patent No. 5869615
; Patent NormATION:
APPLICANT: Dennis E. Hourcade and Teresa J. Oglesby
TITLE OF INVENTION: MODIFIED COMPLEMENT PROTEASES
; NUMBER OF SEQUENCES: 62
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          片
ADDRESSEE PATICAL DATE
ADDRESSEE PATICAL DATE
STREET: 2800 One Atlantic Center
CITY: Atlanta
COUNTRY: USA
ZIP: 30309-3450
COMPUTER: Georgia
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PATICATION DATA:
APPLICATION NUMBER: US/8/177,109A
FILING DATE: 03-UAN-1994
CLASSIFICATION: 514
ATTCHENT APPLICATION NUMBER: WJ/8/177,109A
FILING DATE: 03-UAN-1994
ATTCHENT/ACENT INFORMATION:
REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: WU 107
TELEPOMUNICATION INFORMATION:
TELEPHONE: (404) 873-8794
TELEPHONE: (404) 873-8794
INFORMATION FOR SEQ ID NO: 56:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: mino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMFUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIA Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/482,528
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Guthrie, Janice
REGISTRATION NUMBER: 35,170
REFERENCE/DOCKET NUMBER: 35,170
REFERENCE/DOCKET NUMBER: IT-4630CIP4
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 41.0%;
Best Local Similarity 57.1%;
Matches 4; Conservative 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C11.
STATE: USA
COUNTRY: USA
TTP: 92713-5210
READABLE F
F1'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 25; DB 3; Length 12; Pred. No. 1.3e+02; Indels 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 12;
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Search completed: May 18, 2004, 10:06:44 Job time : 13.75 secs
                                                                                                                                                                                       MOLECULE TYPE: protein; HYPOTHETICAL: NO US-08-177-109A-56
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                                                                                                                            Matches
                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                                   TOPOLOGY:

 Conservative

                                                                                          8 MHNLG 12
                                                               1 LHNMG 5
                                                                                                                                                                                                                                   linear
                                                                                                                         39.3%; Score 24; DB 2; Length 5; 60.0%; Pred. No. 3e+05; tive 2; Mismatches 0; Indels
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Result
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Maximum DB
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Run
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OM protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          on:
number of
                                                                                                                                                                                                                                                                                                                                                                                                                     Pred. No.
                                                                                                                                                                                                                                                                                                                                       Score
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        seq length: 0 seq length: 12
                                                                                                                                                                                                                                                                                                                                                                                              No. is the number of results predicted by chance to have a greater than or equal to the score of the result being printed, s derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-730-174A-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   May 18, 2004, 09:56:10 ; Search time 10.25 Seconds (without alignments)
112.614 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                        Query
Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               283366 seqs, 96191526 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 AVSEIQLMHNLG 12
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                                           Copyright
                                                                                                                                                                                                                                                                                                                                                                                                                                            pir1:
pir2:*
pir3:**
                                                                                                                                                                                                                                                                                                                                        Length
GenCore version (c) 1993 - 2004
                                                                                                                                                                                                                                                                                                                                          DB
                                                                                                                                 S29735
PW0002
A56029
                                                                                                                                                                   146868
A37268
PH1190
PH1187
PH1189
A47364
PH0924
S25266
 E49033
A38841
B41835
B41835
I38888
T09512
PT0238
SPPGNK
S28055
S27873
C61033
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                                                                                                                                            sperm-activating peantigen p20e pre-
5-aminoimidazole r
alpha-myosin heavy
Ig heavy chain C r
T-cell receptor al
T-cell receptor al
T-cell receptor al
placental lactogen
r-cell receptor be
pilE protein - Se
polyphosphate-gluc
chlorophyll a/b-bi
                               COI intron 16 prot
NADH2 dehydrogenas
Ig heavy chain CRD
neuromedin K - pig
cytochrome b559 co
                                                                                                                                                                                                                                                                                                         Ig heavy chain V carnocin UI49 - (
                                                                                       translation elonga
glycoprotein H-a -
                                                                                                                        N-methylpurine DNA
T-cell receptor ga
                                                                                                                                                                                                                                                                                             ribosomal protein
                                                                                                                                                                                                                                                                                                                                           Description
                                                                                                               T-cell receptor ga
rhodopsin homolog
          hypothetical prote ranatachykinin C -
  amine oxidase (cop
                                                                                                                                                                                                                                                                                                                                                                                                                    RyMakiya, R.; Stidbrand, T.

RyMakiya, R.; Stidbrand, T.

Bur. J. Biochem. 205, 341-345, 1992

Bur. J. Biochem. 205, 341-345, 1992

A; Telle: placental alkaline phosphatase has a binding site for the human immunoglobulin A; Telesconce number: S21205; MUID:92209522; PMID:1555592

A; Accession: S21205

A; Accession: S21205

A; Status: preliminary

A; Molecule type: protein

A; Residues: 1-12 < MAK>

C; Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 1
$21205
$21205
Peavy chain V region - human
Ig heavy chain V region - human
C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 16-Aug-1996
C;Accession: $21205
C;Accession: $21205
   RESULT
S78420
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A58718
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Best Local
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35.6%;

; Score 21; DB; ; Pred. No. 6.9et 3; Mismatches

DB 2;

Length 12,

0 Gaps

0

ALIGNMENTS

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R;Stoffels, G.; Nissen-Meyer, J.; Gudmundsdottir, A.; Sletten, K.; Holo, H.; Nes, I.F. Appl. Environ. Microbiol. 58, 1417-1422, 1992
A;Title: Purification and characterization of a new bacteriocin isolated from a Carnoback A;Reference number: A58718; MUID:92321768; PMID:1622206
A;Accession: A58718
A;Status: preliminary
                                                                                                                                                      A;Molecule type: protein
A;Residues: 1-7 <STO>
C;Keywords: antibiotic; lanthionine
                                                                                                                                                                                                                                                                                                                                    C;Species: Carnobacterium sp.
C;Date: 23-Jan-1998 #sequence_revision 30-Jan-1998 #text_change 30-Jan-1998
C;Accession: A58718
                                                                                                                                                                                                                                                                                                                                                                                             carnocin UI49 - Carnobacterium sp. (fragment)
                                                                                               Similarity
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                                    SEIQ 6
SEIQ 5
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100.0%; Er
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; Pred. No. 2.8e+05;
0; Mismatches 0;
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.as 2; Indels
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ribosomal protein RL41, mitochondrial [validated] - rat (fragment) C;Species: Rattus norvegicus (Norway rat) C;Date: 25-Feb-1998 #sequence_revision 13-Mar-1998 #text_change 21-Jul-2000 C;Accession: S78420 K;Gcasck, H.R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Yoshino, K.I.; Kajiura, H.; Nomura, K.; Takao, T.; Shimonishi, Y.; Kurita, M.; Yamagud Comp. Biochem. Physiol. B 94, 739-751, 1989
A;Title: A halogenated amino acid-containing sperm activating peptide and its related peotus nudus, Echinometra mathaei and Heterocentrotus mammillatus.
A;Reference number: A60527
                                                                                                                                                                                                             A;Variety: subtype adr
G;Date: 18-Jun-1993 #sequence_revision 08-Nov-1996 #text_change 15-Aug-1997
C;Accession: B4653
R;Takahashi, K.; Kishimoto, S.; Ohori, K.; Yoshizawa, H.; Machida, A.; Ohnun
J. Immunol. 147, 3156-3160, 1991
A;Title: Molecular heterogeneity of e antigen polypeptides in sera from car:
A;Reference number: A46453; MUID:92013147; PMID:1717588
A;Accession: B46453
                                                                                                                                                                                                                                                                                                                                                                                    e antigen p20e precursor - hepatitis B virus (subtype adr) (fragment) N;Alternate names: HBe antigen precursor N;Contains: e antigen C;Species: hepatitis B virus, HBV
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A;Residues: 1-10 <YOS>
C;Superfamily: unassigned animal
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C;Species: Heterocentrotus mamillatus
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 10-Aug-2000
C;Accession: A60589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Residues: 1-9 <GOL>
A;Note: the protein is designated as mitochondrial ribosomal C;Keywords: mitochondrion; protein biosynthesis; ribosome
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A; Accession: $78420
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A;Residues: 1-10 <TAK>
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                                                                                                                                                                                               A;Status: preliminary
                                                                                    Experimental source: subtype adr;
Note: sequence extracted from NCBI backbone (NCBIP:60243);
1-10/Domain: signal sequence (fragment) #status predicted;
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Similarity
3; Conserv
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75.0%;
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60.0%;
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pred. No. 2.1e+03;
1; Mismatches (
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Pred. No.
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Pred. No. 2.1e+03;
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5-aminoimidazole ribonucleotide carboxylase/4-N-succinylamino carbonyl-5-aminoimidazole C;Species: Gallus gallus (chicken)
C;Date: 19-Gan-1996 #sequence_revision 19-Jan-1996 #text_change 17-Mar-1999
C;Accession: A55837
R;Firestine, S.M.; Davisson, V.J.
Blochemistry 33, 11917-11926, 1994
A;Title: Carboxylases in de Novo purine biosynthesis. Characterization of the Gallus gal A;Reference number: A55837; MUID:95001903; PMID:7918410
A;Accession: A55837
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A55837
                                                                                                                           C;Accession: A37268
R;Ruff-Jamison, S.; Campos-Gonzalez, R.; Glenney Jr., J.R.
J. Biol. Chem. 266, 6607-6613, 1991
A;Title: Heavy and light chain variable region sequences and A;Reference number: A38740; MUID:91177923; PMID:1706720
A;Accession: A37268
                                                                                                                                                                                                                                    Ig heavy chain C region (129) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 19-Mar-1997 #sequence_revision 13-Mar-1998 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;Friedman, D.J.; Umeda, P.K.; Sinha, A.M.; Hsu, H. Proc. Natl. Acad. Sci. U.S.A. 81, 3044-3048, 1984 A;Title: Characterization of genomic clones specifying A;Reference number: I46868; MUID:84221901; PMID:6328491 A;Accession: I46868
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             alpha-myosin heavy chain - rabbit (fragment)
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997
C;Accession: I46868
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A;Molecule type: protein
A;Residues: 1-12 <FIR>
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A; Residues: 1-7 < FRI>
                                                                       A; Molecule type: mRNA
A; Residues: 1-10 < RUF >
                                                                                         A;Status: preliminary A;Molecule type: mRNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Cross-references: GB: K01698; NID: g165538;
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                   28.8%;
Score 17; DB:
Pred. No. 3.3e
1; Mismatches
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Pred. No.
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BIOLWHNIG

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T-cell receptor alpha chain V region (Cw3/2C3) - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Apete: 16-Uul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
C;Accession: PH1189
R;Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; Ba
J. Exp. Med. 176, 439-447, 1992
A;Title: H-2-restricted cytolytic T lymphocytes specific for HLA display T c
A;Reference number: S26512; MUID:92364546; PMID:1380061
A;Accession: PH1189
A;Astatus: preliminary
A;Molecule type: mRNA
A;Residues: 1-12 <CAS>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
C;Accession: PH1187
R;Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; Barra, C.; Wid J. Exp. Med. 176, 439-447, 1992
A;Title: H-2-restricted cytolytic T lymphocytes specific for HLA display T cell receptor A;Reference number: S26512; MUID:92364546; PMID:1380061
A;Accession: PH1187
A;Status: preliminary
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A;Title: H-2-restricted cytolytic T lymphocytes specific for HLA display T cell receptor A;Reference number: S26512; MUID:92364546; PMID:1380061
A;Accession: PH1190
A;Accession: PH1190
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C;Species: Homo sapiens (man)
C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
C;Accession: PH1190
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A;Molecule type: mRNA
A;Residues: 1-12 <CAS>
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C;Species: Homo sapiens (man)
C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
C;Accession: PH1187
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Best Local S
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Best Local S
Matches 4
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                                                        Similarity 100 4; Conservative
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100.0%; Pr
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0;
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0; Mismatches
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                                                             Score 17; DB 2;
Pred. No. 4e+03;
0; Mismatches
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RESULT 12
A47364
A47364
placental lactogen-I precursor - mouse (fragment)
C;Species: Mus sp. (mouse)
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 12-Aug-1996
C;Accession: A47364
R;Shida, M.M.; Ng, Y.K.; Soares, M.J.; Linzer, D.I.
Mol. Endocrinol. 7, 181-188, 1993
A;Title: Trophoblast-specific transcription from the mouse placental lactog A;Reference number: A47364; MUID:93225959; PMID:8469232
A;Accession: A47364
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Maccession: A47364
A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                 pilE protein - Escherichia coli (fragment)
(/Species: Escherichia coli
(/Species: Escherichia coli
(/Species: Escherichia coli
(/Date: 28-May-1993 #sequence_revision 28-May-1993 #text_change
(/Accession: S25266
(/Accession: S2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A, Experimental source: concanavalin A-activated lymphoblast C_1 Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          R;Gold, D.P.; Offner, H.; Sun, D.; Wiley, S.; Vandenbark, A.A.; Wilson, D.B.
J. Exp. Med. 174, 1467-1476, 1991
A;Title: Analysis of T cell receptor beta chains in Lewis rats with experimental allerging A;Reference number: PH0891; MUID:92078857; PMID:1836012
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C; Accession: PH0924
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C;Species: Rattus norvegicus (Norway rat)
C;Date: 09-Oct-1992 #sequence_revision 09-Oct-1992 #text_change 30-May-1997
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A; Residues: 1-10 < RES>
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A; Residues: 1-11 <GOL>
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                                                                                                                                      A;Gene:
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A; Residues: 1-7 < DUP>
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50.0%;
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   Score 15; DB | Pred. No. 2.8e | 1; Mismatches
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Pred. No. 5
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Pred. No. 5.1e+0
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5 IOLM 8
Db 4 IELM 7

RESULT 15
S29735
S29735
C)Apceles: Projoinbacterium freudenreichi subsp. sharmani
C)Species: Projoinbacterium freudenreichi
C)April Dib Horn, PJ.; Wood, H.G.
RESULT 15: PJ.; Wood, H.G.
RESULT 15: P.J.; Wood, H.G.
RESULT 15: Projoinbacterium freudenreichi
C)Species: Projoinbacterium freudenreichi
Result 15: Projoinbacterium freudenreichi
Result 15: Projoinbacterium freudenreichi
C)April 10: PMT 1
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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 12
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Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                             Score
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Match
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Gapop 10.0 , Gapext 0.5
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59
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    ALLAC CARUI
ALLAC CASUP
EPG GLOPA
PVKI PERAM
TRPI PSEPU
ALLI CARMA
ALLI CAPAN
FARB MACRS
UFO MACRS
ALLI CAPAN
TEME RANCA
ASLI BACSE
CD11 LITXA
CON1 PETAM
RS19 TOBB
LCKA LEUMA
ANOSH CLYJA
PORD METIM
RS19 TOBB
LCKA LEUMA
ANOSH CLYJA
PORD
CS15 BACSU
Q20A COMFU
PTBB PAPHA
RE42 LITRU
GLEM HUMAN
RR42 LITRU
GLEM HUMAN
RR42 TOTPA
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p36960 carnobacter p36967 cydia pomon p81010 fusariim so p81350 clostridium p41837 periplaneta p36414 pseudomonas p82158 cydia pomon p81825 cydia pomon p81825 cydia pomon p81826 carcinus ma p82152 cydia pomon p83281 macrobrachi p38640 mus musculu p81822 carcinus ma p56923 rana tempor p22690 rana catesb p01292 sus sorofa p83146 bacteroides p56245 litoria xan p56246 litoria xan p56251 tomato big p21143 leucophaea p19980 leucophaea p19980 leucophaea p19980 periplaneta p35946 phocine dis p810464 comamonas t p82075 homo sapien p81749 clupea pall
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`	P81675 pin			4		P81672 pin	P58807 con					P80678 che
	pinus pinas	itoria rub	riplaneta	cteroides	oteus vui	spurd snurd	conus marmo	rana ridibu	campylobact	heliothis z	romalea mic	chelyosoma

ALIGNMENTS

RESULT 1 LANC CARUI LANC CARUI IDANC CARUI TLANC CARUI DT 01-JUN-1994 (Rel. 2 DT 01-JUN-1994 (Rel. 2 DT 01-JUN-1994 (Rel. 2 DT 16-OCT-2001 (Rel. 4 DE Lantibiotic carnocic Carnobacterium sp. CC Carnobacterium sp. RA Nes I.F., RA SEQUENCE RA MEDLINE=92321768; P. RA CARNOBACTERIUM SP. RA Appl. Environ. Nicr CC -1- PUNCTION. Lanth CC Antibiotic; Bacterium CC -1- PUNCTION. Lanth CC Antibiotic; Bacterium CC -1- PUNCTION. Lanth CC Antibiotic; Bacterium CC -1- FUNCTION. Lanth CC Antibiotic; Bacterium CC Sc Gudaryota; Metazoa; CC Gudaryota; Metazoa; CC Gudaryota; Metazoa; CC Godiastatin 6. CC Godiastatin 6. CC Gudaryota; Metazoa; CC Godiastatin 6. CC Godiastati
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ID_ EFG CLOPA
AC P81350,
DT 15-UUL-1998 (
DT 15-UUL-1998 (
DT 28-FEB-2003 (
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Matches 2
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Verma J., Gangal S.V.;
Submitted (JUL-1997) to Swiss-Prot
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MOD RES 8 8
SEQUENCE 8 AA; 936 MW
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Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordari.
Hypocreomycetidae; Hypocreales; Nectriaceae; Nectria.
NCBI_TaxID=70791;
                                                                                                                                                                           MEDLINE=98291870; PubMed=9629918; Flengsrud R., Skjeldal L.; "Two-dimensional gel electrophoresis separation and N-termin sequence analysis of proteins from Clostridium pasteurianum Electrophoresis 19:802-806(1998).
-!- FUNCTION: This protein promotes the GTP-dependent transl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                P81010;
15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence up
10-OCT-2003 (Rel. 42, Last annotation
Allergen Fus s I3596* (Fragment).
Fusarium solani (subsp. pisi) (Nectria
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                         SUBCELLULAR LOCATION: Cytoplasmic.
SIMILARITY: Belongs to the GTP-binding
EF-G/EF-2 subfamily.
                                                                                                                                                 the nascent protein chain
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01-UUN-1994 (Rel. 29, Last sequence update)
15-MAR-2004 (Rel. 43, Last annotation update)
HTH-type transcriptional regulator trpI (TrpBA
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01-JUN-1994
            Biochimie 71:521-531(1989).
-!- FUNCTION: ACTIVATES THE
                                                                     putida.
                                                                                                  "DNA sequence of the
                                                                                                                                                              MEDLINE=89335826; PubMed=2503057;
                                                                                                                                                                                          SEQUENCE FROM N.A. STRAIN=PPG1 C1S;
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01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last seque
16-OCT-2001 (Rel. 40, Last annot
Periviscerokinin-1 (Pea-PVK-1).
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                                                                                                                 MOD RES
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroidea; Dictyoptera; Blattaria; B
Blattidae; Periplaneta.
NCBI_TaxID=6978;
                                                                                                                                                                         TISSUE=Abdominal perisympathetic organs;

MEDLINE=9523021; PubMed=7716075;

Predel R., Linde D., Rapus J., Vettermann S., Penzli
"Periviscerokinin (Pea-PVK): a novel myotropic neuro
"perisympathetic organs of the American cockroach.";
Peptides 16:61-66(1995).
i- FUNCTION: MYOACTIVE PEPTIDE; HAS EXCITORY ACTION
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InterPro; IPR000847; HTH_LYSR.
PROSITE; PS50931; HTH_LYSR; PARTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cydia pomonella (Codling moth).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neopbera; Endopberygota; Lepidopbera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Duve H., Johnsen A.H., Maestro J.-L., Scott Davey M., East P.D., Thorpe A.; "Lepidopteran peptides of the allatostatin Peptides 18:1301-1309(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Larva;
MEDLINE=98054539; PubMed=9392829;
Duve H., Johnsen A.H., Maestro J.
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30-MAY-2000
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MOD RES 7 7 AMIDATION.
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Pred. No. 1.4e
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Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Bucarida; Decapoda; Pleocyemata; Brachyu
Eubrachyura; Portunoidea; Portunidae; Carcinus.
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|- SIMILARITY: Belongs to the allacostatin family
Neuropeptide, Amidation; Multigene family
MOD RES 8 8 AMIDATION (POTENTIAL)
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
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MOD_RES 8 8
SEQUENCE 8 AA; 934 MW
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-!- SIMILARITY: Belongs to the allatostatin
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TISSUE=Larva;
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P83281;
28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
FMRFamide-like neuropeptide FLP8 (VSHNNFLRF-amide).
Macrobrachium rosenbergii (Giant fresh water prawn)

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MEDLINE-95009907; PubMed=7523108;

Merrick B.A., Patterson R.M., Wichter L.L., He C., Selkirk Merrick B.A., Patterson R.M., Wichter and novel murine programment of the sequencing of familiar and novel murine prousing preparative two-dimensional gel electrophoresis.";

Electrophoresis 15:735-745(1994).

-i- MISCELLANEOUS: On the 2D-gel the determined pI of this
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P38640;
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-i- MASS SPECTROWETRY: MW=1133.8; METHOD=MALDI.
-i- SIMILARITY: Belongs to the FARP (FMRFamide related personal properties of the party 
                                                                                                    -AL19_CARMA
P81822;
30-MAY-2000
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Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Caridea
Palaemonoidea; Palaemonidae; Macrobrachium.
NCBI_TaxID=79674;
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01-OCT-1994 (Rel. 30, Last sequence update)
15-VAR-2004 (Rel. 43, Last annotation update)
Unknown protein from 2D-page of fibroblasts (P32) (Fragment)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Q03367;
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NCBI_TaxID=6759;
               PROSITE; PSOC
Chloroplast;
NON_TER
                                  INDL. ...
PIR; S28055; S250000
HAMAP; MF 00043; -; 1.
Interpro; IPR006216; Cyt b559.
Interpro; PS00537; CYTOCHROME B559;
Interpro; PS00537; CYTOCHROME B559;
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                                                                                                                                                                                                                                  EMBL; X65570; CAA46539.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             between
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2; Conserv
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SEQUENCE.
SEQUENCE.
TISSUB-Cerebral ganglion, and Thoracic
MEDLINE-98121193; PubMed-9461295;
Duve H., Johnsen A.H., Maestro J.-L., S
                                                                                                                                                                                                    STRAIN.cv. Lamuyo; TISSUE-Fruit, and Leaf;
MEDLIKE-93099270, PubMed=1463853;
MEDLIKE-93099270, PubMed=1463853;
Muntz M., Camara B., Weil J.-H., Schantz R.;
The psbL gene from bell pepper (Capsicum annuum): plastid editing also occurs in non-photosynthetic chromoplasts.";
Plant Mol. Biol. 20:1185-1188(1992).
-i- FUNCTION: This b-type cytochrome is tightly associated reaction center of photosystem II and possibly is part water-oxidation complex.
-i- SUBUNIT: Heterodimer of an alpha subunit and a beta sub-
-i- SUBUNIT: Heterodimer of an alpha subunit and a beta sub-
-i- SUBUNIT: Heterodimer of the psbE / psbF family.
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Eubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Viridiplantae; Streptophyta; Embryo
Spermatophyta; Magnoliophyta; eudicotyledons;
lamiids; Solanales; Solanaceae; Capsicum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JUN-1994 (Rel. 29, Created)
01-JUN-1994 (Rel. 29, Last Sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
Cytochrome b559 beta subunit (PSII reaction c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -!- FUNCTION: May act as a neurotransmitter or neu-
-!- SIMILARITY: Belongs to the allatostatin family
Neuropeptide, Amidation, Multigene family.
MOD_RES 10 10 AMIDATION (POTENTIAL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Isolation and identification of multiple neuropeptides allatostatin superfamily in the shore crab Carcinus maen Eur. J. Biochem. 250:727-734(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and foestities requires a license agreement (See http://www.isb-sib.or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Capsicum annuum (Bell pepp
Chloroplast.
Eukaryota; Viridiplantae;
                                                                        s SWISS-PROT entry is copyright. It is produced through ween the Swiss Institute of Bioinformatics and the EW European Bioinformatics Institute. There are no restr by non-profit institutions as long as its content.
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2; Mismatches
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edons; core eudicots; asterids;
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30-MAY-2000
16-OCT-2001
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DOMAIN
SEQUENCE
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P22690;

P22690;

01-AUG-1991 (Rel. 19, Created)

01-AUG-1991 (Rel. 19, Last sequence update)

10-OCT-2003 (Rel. 42, Last annotation update)

Ranatachykinin C (RTK C).

Rana catesbeiana (Bull frog).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Rana.

MCBI TaxID=8400;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TISSUE=Skin secretion;
MEDLINE=97175050; PubMed=9022710;
Simmaco M., Mignogna G., Canofeni S.,
                    TISSUE=Intestine;

MEDLINE=91254337; PubMed=2043143;

MEDLINE=91254337; PubMed=2043143;

Kozawa H., Hino J., Minamino N., Kangawa K., Matsuo H.;

"Isolation of four novel tachykinins from frog (Rana cabrain and intestine.";
                                                                                                                                                                                                                                                                                                                                                                                     -!- SUBCELLULAR LOCATION: Secreted.
-!- TISSUE SPECIFICITY: Skin.
-!- SIMILARITY: Belongs to the brevinin family.
Amphibian defense peptide; Antibiotic; Amidation.
MOD RES 10 10 AMIDATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Rana temporaria (European common frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Ranoidea; Ranidae; Rana.
NCBI_TaxID=8407;
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temporaria."
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          Biochem. Biophys. Res. Commun.
                                                                                 SEQUENCE, AND SYNTHESIS.
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CTION: Has antibacterial activity
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(Rel.
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Pred. No. 7.7e.
4; Mismatches
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Pred. No. 7.7e+03;
1; Mismatches 1;
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LUMENAL (POTENTIAL).
817D0F59D6D69DC5 CRC64;
           177:588-595(1991)
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7.7e+03;
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Best Local :
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TISSUE=Intestine;
MEDLINE=94023216; PubMed=8210506;
MEDLINE=94023216; PubMed=8210506;
Kangawa K., Kozawa H., Hino J., Minamino N., Matsuo H.;
Kangawa K., Kozawa H., Hino J., Minamino N., Matsuo H.;
"Four novel tachykinins in frog (Rana catesbeiana) brain and
intestine.",
Regul. Pept. 46:81-88(1993).
-!- FUNCTION: Tachykinins are active peptides which excite neurons,
evoke behavioral responses, are potent vasodilators and
secretagogues, and contract (directly or indirectly) many smoot!
                                                                                                                                                                        Tachykinin; Neuropeptide; Amidation.
MOD_RES 10 10 AMIDATI
                                                                                                                                                                                                     Interpro; IPR002040; Tachy_Neurokinin.
PROSITE; PS00267; TACHYKININ; 1
                                                                                                                                                                                                                                            -!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: Belongs to the tachykinin family.
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Pred. No.
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seq length: 12
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sp_mammal:*
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sp_bacteria:*
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P82081 PRELIMINARY; PRT; 12 AA.
P82081;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
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100 B 7 S S & W N D .	SCORE 237 237 200 200 200 200 200 118	Match Length 45.8 11 39.0 12 33.9 8 33.9 8 33.9 18 33.9 12 33.9 9 32.2 9 32.2 9 32.2 9 32.2 10		133 133 133 133 133 133 133 133 133 133	Q9PS71 P82081 P82082 P82083 P82083 P82086 Q34909 Q34909 Q92766 Q40659 Q40659	P 10
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ALIGNMENTS

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Q9BS71
Q1-MAY-2000 (TrEMBLrel. 13, Created)
Q1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
Q1-MAY-2000 (TrEMBLrel. 21, Last sequence update)
Q1-JUN-2002 (TrEMBLrel. 21, Last sequence update)
Pibrinolytic metalloproteinase (Fragment)
Agkistrodon contortrix
BUKATYOTA; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea;
Viperidae; Crotalinae; Agkistrodon.
NCBI TaxID=8720;
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NON TER
SEQUENCE
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SEQUENCE.

MEDIINE=91378546; PubMed=1898066;

Guan A.L., Retzios A.D., Henderson G.N., Markland F.S.Jr.;

Guan A.L., Retzios A.D., Henderson of a fibrinolytic enzyme

"Purification and characterization of a fibrinolytic enzyme

"Purification and characterization of a fibrinolytic enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of the southern copperhead snake (Agkistrodon contortrix).";
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ER 11 11
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                                                                               6 QLMHNLG 12
ELGHNLG 10
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71.4%;
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Raftery M.J., Bradford A.M., Bowie J.H., Wallace J.C., Tyler M.J.;
"Peptides from Australian frogs. The structure of the dynastins from the banjo frogs Limnodynastes interioras, Limnodynastes dumerilii a Limnodynastes terraereginae.";
Australia J. Chem. 46:83-842(1993).
-i- MASS SPECTROWETRY: MW=1236; METHOD=FAB.
SEQUENCE 12 AA; 1236 MW; 147AA70FDF472724 CRC64;
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Limnodynastinae; Limnodynastes.
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-!- MASS SPECTROMETRY: MW=772; METHOD=FAB.
SEQUENCE 8 AA; 772 MW; 7B58772455A2C728 CRC64;
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Amphibia; Batrachia; Anura; Neobatrachia;
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NCBI_TaxID=39404;
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a; Craniata; Vertebrata; Euteleostomi;
Neobatrachia; Bufonoidea; Myobatrachi
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Q1-NOV-1996 (TrEMBLrel. 01, C
01-NOV-1996 (TrEMBLrel. 01, I
01-JUN-2003 (TrEMBLrel. 24, I
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TISSUE-SKIN SECRETION, Bowle J.H., TESUE-SKIN SECRETION, Bowle J.H., The structure of the dy Bradford A.M., Raftery M.J., Bowle J.H., The structure of the dy Peptides from Australian frogs. The structure of the dy Peptides from Limnodynastes salmini and Fletcherin from Limnodynastes Limnodynastes salmini and Fletcherin from Limnodynastes Salmini Australy 1244(1993).

Aust. J. Chem. 46:1235-1244(1993).

Aust. J. Chem. 46:1235-1244(1993).

Aust. J. Chem. 46:1235-1244(1993).

Aust. J. Chem. 46:1235-1244(1993).

Aust. J. Chem. 46:1235-1244(1993).
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TISSUB-SKIN SECRETION;

Bradford A.M., Raftery M.J., Bowie J.H., Wallace J.C., Tyler M.J.;

"Peptides from Australian frogs. The structure of the dynastins from

Limnodynastes salmini and Fletcherin from Limnodynastes fletcheri.";

Aust. J. Chem. 46:1235-1244(1993).

-!- MASS SPECTROMETRY: MW=1114, METHOD=FAB.

SEQUENCE 12 AA; 1114 MW; 3AB5A976CAA72728 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Limnodynastes salmini (Salmin's-striped frog).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Ev
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea;
Limnodynastinae; Limnodynastes.
SEQUENCE FROM N.A.

MEDLINE=8823478; PubMed=2836084;

McCracken A., Uhleabusch I., Gellissen G.;

"Structure of the cloned Locusta migratoria mitochondrial

"Struction mapping and sequence of its ND-1(URF-1) gene.

Curr. Go.; GO:0005739; C:mitochondrion; IEA.
                                                                                                                                                                                                                                      Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroidea; Orthoptera; Caelifera; Av
Acridoidea; Acrididae; Oedipodinae; Locusta.
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Tynastins from
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SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

C STRAIN=Dog #10757/96;

A Liermann H., Harder T., Haas L.;

A Liermann H. Harder T., Haas L.;

T "Genetic analysis of the central untranslated genome reg

RT proximal coding part of the F gene of wild-type and vacc

RT morbilliviruses.";

RL Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases

FMRI.; AF026234; AAC09164.1; -.
              STREET REAL STREET
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STRAIN=DOG #5526/89;
Liermann H., Harder T., Haas L.;
Liermann H., Harder T., Haas L.;
"Genetic analysis of the central untranslated genor proximal coding part of the F gene of wild-type and proximal coding part of the EMBL/GenBank/DDBJ data morbilliviruses.";

Submitted (SEP-1997) to the EMBL/GenBank/DDBJ data EMBL, AF026237; AAC09167.1; -.

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SEOÜENCE 9 AA; 1011 MW; F281732760533441 CRC64
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01-AUG-1998 (TrEMBLrel. 07, C1
01-AUG-1998 (TrEMBLrel. 07, L.
01-DEC-2001 (TrEMBLrel. 19, L.
Fusion protein (Fragment).
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Viruses; ssRNA negative-strand viruses; Mononegavirales;
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Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.
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pred. No. 1e+06;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
Alpha-amylase (Fragment)
Oryza sativa (Rice).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
NCBI TaxID=4530;
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OPEXC3;
OPEXC3;
O1-MAY-2000 (TrEMBLrel. 13, Created)
O1-MAY-2000 (TrEMBLrel. 13, Last sequence update)
O1-JUN-2003 (TrEMBLrel. 24, Last annotation update)
E antigen P20E (Fragment).
Hepatitis B virus.
Viruses; Retroid viruses; Hepadnaviridae; Orthohepadnavirus.
NCBI_TaxID=10407;
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EMBL; M62916; AAA3389
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MEDLINE=91078641;
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Takahashi K., Kishimoto S., Ohori K., Yoshizawa H., Ma Ohnuma H., Tsuda F., Munekata E., Miyakawa Y., Mayumi "Molecular heterogeneity of e antigen polypeptides in carriers of hepatitis B virus.";
J. Immunol. 147:3156-3160(1991).
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Q28742 PRELIMINARY; PRT; 7 AA. Q28742; Q28742; Q1.NOV-1996 (TrEMBLrel. 01, Created) Q1.NOV-1996 (TrEMBLrel. 01, Last sequence update) Q1.JUN-2003 (TrEMBLrel. 24, Last annotation update) Alpha-myosin heavy chain (Fragment). Qryctolagus cuniculus (Rabbit).
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01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
MH2, proviral DNA, myc to 3' LTR (Fragment).
Avian carcinoma virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=82260445; PubMed=6286214; MCKecwn M., Firtel R.A.; "Actin multigene family of Dictyostelium. Cold Spring Harb. Symp. Quant. Biol. 46:4 EMBL; K02957; AAA33150.1; -. EMBL; K02956; AAA33150.1; JOINED. SEQUENCE 11 AA; 1205 MW; 728B4C14C6C2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Viruses; Retroid viruses; Retroviridae; Alpharetrovirus
NCBI_TaxID=11958;
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NCBI_TaxID=44689;
[1]
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ID Q9GD1
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C9GD12;
PKBLACC...
Q9GD12;
01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-UN-2003 (TrEMBLrel. 24, Last annotation update)
Ribosomal protein S16 (Fragment).
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01-NOV-1996 (TrEMBLre
01-NOV-1996 (TrEMBLre
01-DEC-2001 (TrEMBLre
(Clone XP2E8B) (Fragm
Homo sapiens (Human).
Chloroplast.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Arecaceae;
                                                                                                                                                                                                                                                                                                                                                                                         "Isolation of chromosome-specific genes arrayed cDNAs and cosmid libraries,", Hum. Mol. Genet. 0.0-0(1995).
EMBL; L32131; AAA73881.1; -.
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Coolbaugh M.I., Chinault C.A.,
Caskey C.T.H.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ventricular myosin heavy proc. Natl. Acad. Sci. U.S
EMBL; K01698; AAA31415.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
MEDLINE=84221901; PubMed=6328491;
Friedman D.J., Umeda P.K., Sinha
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Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TISSUE=Placenta;
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3; Conserv
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4; Conserv
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eavy chains.":
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44.4%;
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U.S.A. 81:3044-3048(1984).
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Last sequence update)
Last annotation update)
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2; Mismatches
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Pred. No. 1e+06;
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Baldini A.,
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, Lindsay E.A., Zhao
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OC Areceae; Linospadicinae; Linospadix.

OX NCBI_TAXID=131282;

RN [1]

RN [1]

RP SEQUENCE FROM N.A.

RC TISSUE=Leaf;

RA ASMUSSEN C.B., Chase M.W.;

RT "Coding and noncoding plastid DNA in palm systematics.";

RI Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.

BR "Coding and noncoding plastid DNA in palm systematics.";

RI Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.

DR GO:0009507; C:chloroplast; IEA.

WW Chloroplast.

BR GO:GO:0009507; C:chloroplast; IEA.

WW Chloroplast.

FT NON_TER 9 9

SQ SEQUENCE 9 AA; 1106 MW; 1B9236C2C0441B50 CRC64;

Query Match 28.8%; Score 17; DB 8; Length 9;

Best Local Similarity 75.0%; Pred. No. 1e+06;

Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Matches 3; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Search completed: May 18, 2004, 10:04:48

Job time : 31.25 secs
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Result
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(KRON/) KRONENBERG H M.
(POTT/) POTTS J T.
(JUEP/) JUEPPNER H.
                                                                                                                                                                                                                    sapiens.
                                                                                                                                                                                                                                parathyroid hormone; N-terminal; signaling domain; bone marreformation; resorption; remodeling; tether1; osteoporosis
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Compounds of the structure or formula S-(L)_n-B, R_1-S-(L)_n-R or S-(L)_n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH), L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4)), B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-rP) (1-34); R_1 is the PTH-1 receptor signal sequence, and R is (a portion of) PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAMP Claim 4; Page 92; 119pp; English. New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass. for

WPI; 2000-452384/39.

Gardella TJ,

Kronenberg

MH,

Potts JT,

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72607
The invention relates to a new antigenic peptide for inducing the formation and isolation of antibodies having an affinity to it, being formed from the N-terminus of parathyroid hormone (PTH). Also included are; (1) a method for producing antibodies useful in the determination PTH levels in a biological sample comprising: (a) providing at least on first peptide antigen comprising a peptide fragment of PTH; (b) administering the first peptide antigen to a host animal to induce antibody production; (c) monitoring the antibody titre produced; (d) isolating antisera produced in the host animal; and (e) selecting antisera from the isolated antisera produced in the host that is capable of binding to a second peptide antigen; (2) an antibody (ab) produced be
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90.9%;
                                                                                                                                                                                                                               JR;
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Pred. No. 0.012;
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Claim 2;

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New parathyroid hormone (PTH) antigenic peptide inducing the and isolation of antibodies having an affinity to it, useful determining bloactive PTH levels in serum, plasma and/or cell

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Matches 10
                                                                                New antigenic peptide(s) from generated using them, able to forms of the hormone.
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                             Claim
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                             2;
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                                                                                                                                                                                                                                                                                                              FORSSMANN W.
                             Page 4;
                                                                                                                                                                                                                                                     Z,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    thyroid hormone; hPTH; active hPTH 1-37.
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                             5pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hormone antigenic peptide hPTH 1-10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          peptide; 10
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90.9%;
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Pred. No.
                                                                                                           human parathyroid hormone - and antibodies distinguish between active and inactive
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                                                                                                                                                                                                                                                     Maegerlein
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0.014;
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                                                                                                                                                                                                                                    The present sequence represents a parathyroid hormone (pTH) fragment, comprising amino acids 1-10. Parathyroid hormone fragments of the invention have lipolysis stimulating activity (especially when topically administered). The lipolytic activity of the peptides is enhanced when they are chemically modified to increase their lipolyhilicity. The peptides are used in cosmetic or dermatological compositions for skin care. They are especially used for slimming treatment of excessive weight in the thighs and hips, in the treatment of cellulite and for skin firming.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Page 8; 18pp; French
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New parathyroid hormone fragment peptides, in topically applied cosmetic compositions excessive weight in hips and thighs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2000-171243/15.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-JUL-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Parathyroid hormone; lipolysis; cosmetic; de slimming treatment; cellulite; skin firming.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amino acids 1-10 of a parathyroid hormone (pTH)
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                                                                                                                                                                   Sequence 10
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AVSEIQLMHN 10
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                                                              Conservative
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90.0%;
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Pred. No. 0.06
1; Mismatches
                                                            Score 46; DB |
Pred. No. 0.06
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                                                                                                                  DB 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        used as lipolysis stimulants for slimming treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ν.
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                                                                                                                                                                                                                                                                                  This invention describes a novel method for determining (M1) the content of active parathyroid hormone (A) by treating a sample with (i) antibody (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), and including the N-terminal residue and (ii) antibody (Ab2) that creognizes an epitope within the receptor-binding site of (A). The number of molecules that react with both antibodies is determined and used to calculate the content of physiologically active (A). The method is used for diagnosis of calcium-metabolism disorders, osteopathy, or hypo- or contemporare thyroidism. The method (unlike known assays) recognizes that the complete (B4 aa) peptide are content of (A) shorter than the complete (B4 aa) peptide are active and (ii) that apparently intact peptide may be biologically cative, and also takes into account the fact that some fragments of (A) care antisponistic (these have the receptor-binding site but lack the N-complicative (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This sequence represents a peptide fragment used to illustrate the method of
                                                                                                                                                                               Query Match
Best Local S
Matches 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibodies reactive with different epitopes.
                                                                                                                                                                                                                                          Sequence 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Disclosure; Page 3; 10pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2001-376318/40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           hypo-parathyroidism; hyper-parathyroidism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human parathyroid
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                                             ABR44166 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (IMMU-) IMMUNDIAGNOSTIK AG
                                                                                                                                                                                                                                                                            invention
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9; Conserv
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                                                                                                                                                                                  Conservative
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Chang M;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a fusion peptide (Tat-hPTHDP), where self cell-penetrating Tat peptide is bound to human parathyroid hormone-derived peptide (hPTHDP). The fusion peptide is useful as a component of skin slimming cosmecic composition. The fusion peptide does not cause irritation, easily and safely penetrates into integument and endothelium does not cause skin disease and has superior lipplysis effects, and is durable. The present sequence represents a human parathyroid hormone (hPTH) fragment that can be used to construct the fusion peptide
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                                                                                                                                                                                                                                                                                                                                                                                                         Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cyclic peptide; resin bound; parathyroid hormone; osteopathic; disease; hypotensive; PTH receptor; treatment; hyper-calcemia; hypo-calcemia; osteoporosis; osteopenia; hyper-parathyroidism; hypo-parathyroidism; Cushing's syndrome; renal failure; hypertension; bone fracture repair.
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100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel method for the preparation of peptides CC (II) that contain both cyclic and linear peptide fragments comprises CC sequential reaction of a resin-bound linear fragment with the cyclic CC fragment in N-protected form and optionally other linear fragments. The CC products of the invention have osteopathic or antagonists of hPTH. The CC method is particularly used to prepare cyclic peptide analogs of CC granthyroid hormone (PTH) or PTH-related peptides which are useful for CC treating diseases that respond to treatment with agents that bind to PTH CC receptors (with or without activation of adenyl) cyclase activity), e.g. CC hyper- or hypo-calcemia, osteoporosis, osteopenia, hyper- or hypo-calcemia, osteoporosis, osteopenia, hyper- or hypo-calcemia fractures. Separate synthesis of the cyclic for promoting repair of bone fractures. Separate synthesis of the cyclic fragment allows convergent synthesis of resin-bound (II), with better CC preparation of the bridged fragment are confined to a small peptide which can be purified before reaction with the resin-bound component. AAY5568-cc invention
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                                                                                                                                                                                                                                                                          Parathyroid hormone peptide; PTH; PTH-related peptide; PTHrP; calcium homeostasis; PTH-1 receptor; PTH-2; vitamin D synthes bone synthesis; agonist; osteoporosis; non-parenteral deliver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (RHON ) RHONE-POULENC RORER PHARM
                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                        PTH(1-14)/PTHrP(1-14)-derived peptide, SEQ ID NO:6.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB01862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB01862 standard; peptide; 9 AA
                                                                                                                             WO200023594-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1999-633822/54.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AVSEIQLMHNLG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AVSEIQL-HNLG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99WO-US008435
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            73.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 43.5; DB Pred. No. 0.2; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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20-OCT-1999;

99WO-US024481

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RESULT 11
AAY96966
ID AAY96
XX
AC AAY96
XX
DT 31-OC
XX
XX
XX
DE Parat
XX
KW PTH;
                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to a novel parathyroid hormone (PTH) peptide (CAAB01859) and parathyroid hormone related peptide (PTH:P; AAB01861), and biologically active derivatives thereof (AAB01857-801858). AAB01861-CC B01869). The peptides of the invention are at least 85% identical to the CG generic peptide of the formula: X1-Val-Ser-Glu-X2-Gln-Leu-X3-His-X4-X5-CC Gly-Lys-X6 (AAB01857) where: X1 is Ser or Ala; X2 is Ile or Lys; X3 is GG Hy-Lys-X6 (AAB01857) where: X1 is Ser or Ala; X2 is Ile or Lys; X3 is GG Hy-Lys-X6 (AAB01857) where: X1 is Ser or Ala; X2 is Ile or Lys; X3 is GG Hy-Lys-X6 (AAB01857) where: X1 is Ser or Ala; X2 is Ile or Lys; X3 is GG Hy-Lys-X6 (AAB01857) where: X1 is Ser or Ala; X2 is Ile or Lys; X3 is GG Hy-Lys-X6 (AAB01857) where X1 is Ser or Ala; X2 is Ile or Lys; X3 is GG Hy-Lys-X6 (AAB01857) where X1 is Ser or Ala; X2 is Ile or Lys; X3 is GG Hy-Lys-X6 (AAB01857) where X1 is Ser or Ala; X2 is Ile or Lys; X3 is GG Hy-Lys-X6 (AAB01857) where X1 is Ser or Ala; X2 is Ile or Lys; X3 is GG Hy-Lys-X6 (AAB01857) where X1 is Aabout Ala; X2 is Ile or Lys; X3 is GG Hy-Lys-X6 (AAB01857) where X1 is Aabout Ala; X2 is Ile or Lys; X3 is CC Cand is necessary for the normal function of the gastrointestinal, and C-CC entropy of anticy and Aabout Aab
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Best Local
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(POTT/)
(JUEP/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Parathyroid hormone (PTH)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-339693/29
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PTH; parathyroid hormone; N-terminal; signaling
                                                                        31-OCT-2000
                                                                                                           AAY96966;
                                                                                                                                              AAY96966 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 9 AA;
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KRONENBERG H M.
POTTS J T.
                                                                                                                                                                                                                                                                                                              similarity
9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       POTTS J T.
JUEPPNER H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IJ,
                                                                                                                                                                                                                                                                             AVSEIQLMH
                                                                                                                                                                                                                                        AVSEIQLMH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        d hormone (PTH) peptides, PTH related peptides encode them, useful for treating osteoporosis.
                                                                                                                                                                                                                                                                                                                Conservative
                                     hormone N-terminal signaling domain
                                                                        (first entry)
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                                                                                                                                                                                                                                                                             9
                                                                                                                                                                                                                                                                                                      72.9%; 5c.
100.0%; Prr
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                                                                                                                                                                                                                                                                                                                                Score 43;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PTH related peptides and the nucleic
                                                                                                                                                                                                                                                                                                        DB 3; Le..
J. 1.4e+06;
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                                                                                                                                                                                                                                                                                                                                                   Length 9;
   domain; bone mass,
                                         (residues 1-9).
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     28-SEP-1994;
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RESULT 12
AAR91645
ID AAR91
XX AAR91
XX AAR91
XX Humar
XX
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Compounds of the structure or formula S-(L)_n-B, R_1-S-(L)_n-R or S-(L)_n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4)); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-TP) (1-34); R_1 is the PTH-1 receptor signal sequence; and R is (a portion of PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAMP in a mammalian cell having PTH-1 receptors, or screening for a peptide or non-peptide PTH (claimed). The new compound can be administered by inhalation unlike the large native PTH or PTH-PTH which avoids the need for regular injections to treat osteoporosis
                                                                                                                                                                                                                          Human parathyroid hormone; hPTH; diagnosis; active hPTH 1-37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.
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                                                                                                                DE4434551-A1
                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                 Human parathyroid hormone antigenic peptide hPTH 1-9
                                                                                                                                                                                                                                                                                                                                                                                06-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR91645 standard; peptide; 9 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; Page 92; 119pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-452384/39
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(KRON/) KRONENBERG
(POTT/) POTTS J T.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Similarity 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JUEPPNER H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                KRONENBERG H M. POTTS J T.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                (first entry
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100.0%; Pr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 43;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                            antigen; alpha-helix;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.4
DB
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                                                                                                                                                                                                                                                                      immunogen;
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94DE-04434551.

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Best Local S
Matches 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New antigenic peptide(s) from generated using them, able to
                                        Parathyroid hormone (PTH) acids that encode them, us
                                                                                            Gardella TJ,
                                                                                                                           (GARD/)
(KRON/)
(POTT/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 2; Page 4; 5pp; German.
                   Disclosure; Page 26;
                                                                                                                                                                       22-OCT-1998;
                                                                                                                                                                                          20-OCT-1999;
                                                                                                                                                                                                                                     WO200023594-A1
                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                   Parathyroid hormone peptide; PTH; PTH-related peptide; PTHrP; calcium homeostasis; PTH-1 receptor; PTH-2; vitamin D synthes
                                                                                                                                                                                                                                                                                                                                    PTH(1-14)/PTHrP(1-14)-derived peptide, SEQ ID NO:7.
                                                                                                                                                                                                                                                                                                                                                         11-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                               AAB01863
                                                                                                                                                                                                                                                                                                                                                                                                   AAB01863 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                determination of biologically active hPTH(1-37)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Adermann K,
                                                                                                                                                                                                                                                                                          bone synthesis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1996-180391/19.
                                                                        2000-339693/29
                                                                                                                 ) GARDELLA T J.
) KRONENBERG H M.
) POTTS J T.
) JUEPPNER H.
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                             Kronenberg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            94DE-04434551
                                                                                                                                                                       98US-0105530P
                                                                                                                                                                                           99WO-US024481.
                                                                                                                                                                                                                                                                                          agonist;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           67.8%;
                  73pp; English
                                        H) peptides,
useful for
                                                                                             ,
MH
                                                                                                                                                                                                                                                                                          osteoporosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 40; DB:
Pred. No. 1.4e.
1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          distinguish between active and inactive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       human parathyroid hormone - and antibodies
                                                                                              Potts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 'n
                                                                                            JT,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Maegerlein
                                        PTH related peptides and the nucleic reating osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Db 2,
1.4e+06;
0;
                                                                                                                                                                                                                                                                                          TH-2; vitamin D synthesis;
non-parenteral delivery.
                                                                                            Jueppner
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
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The invention relates to

a novel parathyroid hormone

(PTH) peptide

Gardella TJ,

Kronenberg

M M M

Potts JT,

Jueppner H;

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RESULT 14
AAY96981
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CC biologically active derivatives thereof (AABD1857-B01886). The peptides of the invention are at least 85 identical to the CC generic peptide of the formula: XI Val-Ser-Glu-X2-Gln-Leu-X3-His-X4-X5-CC Gly-Lys-X6 (AABD1857) where: X1 is Ser or Ala; X2 is Ile or Lys; X3 is CC Met, Leu or N1e; X4 is Asn or Asp; X5 is Leu or Lys; X6 is His or Ser; CC provided that the peptide is not PTHTP(1-14). The peptides of the CC invention also encompass fragments of peptides of the invention consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C-C terminal derivatives. PTH is a major regulator of calcium homeostasis, CC and is necessary for the normal function of the gastrointestinal, consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C-C terminal derivatives. PTH is a major regulator of calcium homeostasis, CC and is necessary for the normal function of the gastrointestinal consisting to both PTH-1 receptors on osteoblasts and renal tubular cells, CC in the tell to both PTH-1 receptors on osteoblasts and renal tubular cells, CC and to the recently identified PTH-2 receptor. PTH has a potent anabolic calls of protein bridge and vitamin D synthesis in the kidney. A homologous calls of protein agonists of PTH, and also bind to the PTH-1 minic some of the creation are either agonists of PTH, and also bind to the PTH-1 minic some of the conditions characterised by a decrease in bone mass, such as cottened and conditions characterised by a decrease in bone mass, such as cottened conditions characterised by a decrease in bone mass, such as cottened conditions characterised by a decrease in bone mass, such as conditions characterised by a decrease in bone mass, such as conditions characterised by a decrease in bone mass, such as conditions characterised by a decrease in bone mass, such as conditions characterised by a decrease in bone mass, such as conditions characterised by a decrease in bone condition are also conditions characterised by a decrease in bone condition are also conditions character
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Best Local
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                                                                  (GARD/) GARDELLA T J.
(KRON/) KRONENBERG H M.
(POTT/) POTTS J T.
                                                                                                                                                                     31-DEC-1998;
                                                                                                                                                                                                                                                                     06-JUL-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    non-parenteral route:
PTH-1/PTH-2 receptor
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8; Conserv
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88.9%;
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Pred. No.
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1.4e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.
This invention describes a novel of active parathyroid hormone (A)
                                                                                                  Determining the content of physiologically active parathyroid hormone useful in diagnosis of calcium-metabolism disorders, using two antiboo
                                                                                                                                                                                                                                                                                                                                                                                                                      Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist; hypo-parathyroidism; hyper-parathyroidism.
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                                                  Disclosure; Page 3; 10pp;
                                                                                                                                                      WPI; 2001-376318/40.
                                                                                                                                                                                                                                                       17-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human parathyroid hormone immunogenic peptide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAB86220,
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88.9%;
                                                     German
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   /el method for determining (M1) the content
(A) by treating a sample with (i) antibody
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CC (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), CC and including the N-terminal residue and (ii) antibody (Ab2) that CC recognizes an epitope within the receptor-binding site of (A). The number CC of molecules that react with both antibodies is determined and used to CC calculate the content of physiologically active (A). The method is used CC for diagnosis of calcum-metabolism disorders, osteopathy, or hyporocC for parathyroidism. The method (unlike known assays) recognizes that CC (i) some fragments of (A) shorter than the complete (B4 aa) peptide are CC active and (ii) that apparently intact peptide may be biologically cCC inactive, and also takes into account the fact that some fragments of (A) CC are antagonistic (these have the receptor-binding site but lack the N-CC terminus). It thus provides a true measure of the content of CC and its 1-37 fragment which may produce falsely high values. This CC sequence represents a peptide fragment used to illustrate the method of CC the invention.
                                                           the invention
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В Ş 1 AVSEIQLMH SVSEIQLMH φ ω

Matches Query Match Best Local 9

Local Similarity nes 8; Conserv

Conservative

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Score 40; DB 4; Li Pred. No. 1.4e+06; l; Mismatches 0;

Length 9;

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Gaps

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67.8%;

Sequence 9 AA;

Search completed: May 18, 2004, 10:01:54 Job time : 45.75 secs

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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100.425 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

1: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Score

Query Match Length

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Description

RESULT 2 US-09-730-174A-3

Sequence 3, Application US/09730174A Patent No. US20020110871A1

GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a
TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Ho
FILE REFERENCE: IMUNE-001A

ALIGNMENTS

RESULT 1 US-09-730-174A-5

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Sequence 5, Application US/09730174A
Patent No. US20020110871A1
GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Hc TITLE OF INVENTION: MUMBER: US/09/730,174A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 5
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RESULT 3
US-09-730-174A-1
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CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 3
LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                    TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid FILE REFRERICE: INUNE-001A CURRENT APPLICATION NUMBER: US/09/730,174A CURRENT FILING DATE: 2000-12-05 NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J
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SEQ ID NO 1
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TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having
TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid;
FILE REFERENCE: INUNE-001A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
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TYPE: PRT
                                                                                                      ORGANISM: Artificial Sequence
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                                                                            OTHER INFORMATION: Induces Formation of Antibodies and Isolates
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Pred. No. 0.
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US-09-730-174A-4
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US-09-730-174A-2
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Best Local
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TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Hole Fill Reference: IMUNE-001A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 4
LENGTH. 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Induces Formation of Antibodies and US-09-730-174A-4
                                                                                            ; Sequence 1, Application US/10168185; Publication No. US20030175802A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid How FILE REFERENCE: IMUNE.001A CURRENT APPLICATION NUMBER: US/09/730,174A CURRENT FILING DATE: 2000-12-05 NUMBER OF SEQ ID NOS: 12

SEQ ID NO 2
LENGTH: 11
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2, Application US/09730174A Patent No. US20020110871A1
                    APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 12
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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OTHER INFORMATION:
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10; Conserv
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90.9%;
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Pred. No. 0.01
0; Mismatches
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Pred. No.
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Parathormone

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APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 7
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US-10-168-185-7
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; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-168-185-1
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Best Local Similarity
Yatches 9; Conserve
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US-10-192-673-6
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US-10-168-185-7
                                                                                                                                                                Sequence 6, Application US/10192673
Publication No. US20030166838A1
GENERAL INFORMATION:
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                    APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 9
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nes 9; Conserv
REFERENCE: 0609.4570002
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100.0%;
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Pred. No. 0.11
1; Mismatches
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      OF INVENTION:
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APPLICANT: ANDRUSSER, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SECTIONARE: FastSEQ for Windows Version 4.0
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US-10-192-673-7
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Publication No. US20030175802A1
GENERAL INFORMATION:
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Best Local S
Matches 9
                                                           Sequence 7, Application US/10192673
Publication No. US2003016838A1
GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Duppner, Harald
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CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: US. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: US. 60/105,530
PRIOR FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
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TYPE: PRT
APPLICANT: Juppner, Harald TITLE OF INVENTION: Bloactive Peptides and Peptide Derivatives TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: peptide
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                                                                                                                                                                                                                                                                                                                                                                                             Local
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8; Conserva
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100.0%; Pred. No.
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. 1e+06;
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1e+06;
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APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT APPLICATION NUMBER: PCTEP00/12911
PRIOR APPLICATION NUMBER: PCTEP00/12911
PRIOR APPLICATION NUMBER: PCTEP00/12911
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US-10-192-673-8
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CURRENT APPLICATION NUMBER: US/10/192,673

CURRENT FILING DATE: 2002-07-11

PRIOR APPLICATION NUMBER: U.S. 09/421,379

PRIOR FILING DATE: 1999-10-20

PRIOR APPLICATION NUMBER: U.S. 60/105,530

PRIOR FILING DATE: 1998-10-22

NUMBER OF SEQ ID NOS: 13

SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                ; ORGANISM: Homo sapiens
US-10-168-185-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 2, Application US/10168185 Publication No. US20030175802A1 GENERAL INFORMATION:
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                                                                                                              Sequence 8, Application US/10192673
Publication No. US20030166838A1
GENERAL INFORMATION:
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Best Local &
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                                                                                                                                                                                                                                                                                                                                     Matches
                                     APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
    TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 9
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRITY ORGANISM: Artificial Sequence FEATURE:
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CANT: Juppner, Harald
OF INVENTION: Bioactive Peptides and Peptide Derivatives
OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
                                                                                                                                                                                                                                                                                             1 AVSEIQLMH 9
                                                                                                                                                                                                                                                                                                                                     Similarity
8; Conserv
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Pred. No. 1e+06;
1; Mismatches C
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Pred. No. le+06;
1; Mismatches 0
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Sequence 10, Applicate publication No. US200 GENERAL INFORMATION:

Application US/10192673

APPLICANT: Gardella, Thomas J. APPLICANT: Kronenberg, Henry APPLICANT: Potts, John T. APPLICANT: Juppner, Harald

PPLICANT: Juppner, Harald ITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives

```
APPLICANT: Roth, Heinz-Jurgen

APPLICANT: Roth, Heinz-Jurgen

TITLE OF INVENTION: Method for Determining Parathormone

TITLE OF INVENTION: Activity in a Human Sample

FILE REFERENCE: HLZ-004US

CURRENT APPLICATION NUMBER: US/10/168,185

CURRENT FILING DATE: 2002-06-17

PRIOR APPLICATION NUMBER: CCT/EP00/12911

PRIOR FILING DATE: 2000-12-18

PRIOR FILING DATE: 2000-12-18

PRIOR FILING DATE: 1999-12-17

NUMBER OF SEQ ID NOS: 11

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 3

LENGTH: 8

TYPE: PRT

CRGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
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; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide
US-10-192-673-8
RESULT 15
US-10-192-673-10
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Best Local S
Matches 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinricl
APPLICANT: Roth, Heinz-Jurgen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
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7; Conserv
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77.8%;
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Pred. No.
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Pred. No.
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TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid;
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)

FILE REFERENCE: 0609.4570002

CURRENT REPLICATION NUMBER: US/10/192,673;
CURRENT REPLICATION NUMBER: US. 09/421,379
PRIOR APPLICATION NUMBER: US. 09/421,379
PRIOR APPLICATION NUMBER: US. 60/105,530
PRIOR FILLING DATE: 1999-10-20
PRIOR FILLING DATE: 1999-10-22

NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 10
LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE: OTHER INFORMATION: Description of Artificial Sequence: synthetic
OTHER INFORMATION: peptide
US-10-192-673-10

Query Match
Best Local Similarity 66.7%; Pred. No. 1e+06;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps

OY 1 AVSEIQLMH 9

Db 1 SVSEHQLLH 9

Search completed: May 18, 2004, 10:21:00

Search completed: May 18, 2004, 10:21:00
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GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.

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Result
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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length: 12
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| cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
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US-09-421-379-6

US-08-817-547A-2

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US-09-421-379-10

US-08-817-547A-4

US-08-817-547A-4

US-08-817-7109A-56

US-08-817-7109A-56

US-08-817-7109A-56

US-08-817-747A-11

US-08-817-547A-16

US-08-817-547A-16

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US-08-817-547A-5

US-08-817-547A-5

US-08-817-547A-5

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ence 184	e 41,	e 89,	e 27,	Sequence 56, Appl	Sequence 27, Appl	nce 71,	Sequence 73, Appl	e 71,	nce 177		Sequence 10, Appl		e 74,	e 271	Sequence 42, Appl	-	Sequence 6, Appli	

APPLICANT: MENCE!, James J. TITLE OF INVENTION: PROCESS FOR THE PREPARATION OF RESIN-BOUND CYCLIC TITLE OF INVENTION: PROCESS FOR THE PREPARATION OF RESIN-BOUND CYCLIC TITLE OF INVENTION: PAPEL SOFTIDES CURRENT FILLING LATE: 1999-11-18 CURRENT FILLING DATE: 1999-11-18 EARLIER APPLICATION NUMBER: 60/081,897 EARLIER APPLICATION NUMBER: 60/081,897 EARLIER FILLING DATE: 1998-04-15 NUMBER OF SEQ ID NOS: 46 SOFTWARE: PATENTING DATE: 1998-04-15 SOFTWARE: PATENTING DATE: 1998-04-15 SOFTWARE: PATENTING DATE: 1998-04-15 NAME/KEY: PEPTIDE LOCATION: (1) OTHER INFORMATION: FMOC-Ala FEATURE: NAME/KEY: PEPTIDE LOCATION: (3) OTHER INFORMATION: Sex (OtBu) FEATURE: NAME/KEY: PEPTIDE LOCATION: (4) OTHER INFORMATION: Glu (OtBu) FEATURE: NAME/KEY: PEPTIDE LOCATION: (6) OTHER INFORMATION: Glu (OtBu) FEATURE: NAME/KEY: PEPTIDE LOCATION: (6) OTHER INFORMATION: Glu (OtBu) FEATURE: NAME/KEY: PEPTIDE LOCATION: (6) OTHER INFORMATION: Glu (OtBu) FEATURE: NAME/KEY: PEPTIDE LOCATION: (6) OTHER INFORMATION: His (Trt) FEATURE: NAME/KEY: PEPTIDE LOCATION: (9) OTHER PROFRAMATION: His (Trt)
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Sequence 6, Application US/09421379
Patent No. 6495662
GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
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Matches 11
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Best Local Similarity 90.0
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Patent No. 6030790
                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/
PILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 0782
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
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7: USA
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191 Peachtree Street, 37th Floor
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90.0%;
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Pred. No. 0.00:
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Pred. No. 0.019;
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Best Local
                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95,
PILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 078:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-818-3700
                                                                                                                     TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                 MOLECULE TYPE: HYPOTHETICAL:
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STREET: 10-
CHORGIA
CHORGIA
                                                                         TYPE: amino acid
STRANDEDNESS: unl
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                                                                                                       ENGTH:
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9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                           Georgia
Y: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AVSEIQLMH 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AVSEIQLMH 9
                                                                                                          9 amino acids
                                                                                                                                                  404-818-3799
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Adermann, Knut
                                                               unknown
                             .. peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jones & Askew, LLP
                                                                           unknown
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; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-09-421-379-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: U.S. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid TITLE OF INVENTION: Hormone-Related Peptide (PTHrP) FILE REFERENCE: 0609.4570001
                                                                                                                                                                                 SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/817,547A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
NUMBER OF SEQUENCES: 36
                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                               191 Peachtree Street, 37th Floor
                                                                                                                PCT/EP95/03757
07826-0007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 43;
Pred. No.
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3e+05;
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Matches Query Match Best Local

Similarity 8; Conserv

Conservative

67.8%; 88.9%;

1 AVSEIQLMH 9

US-09-421-379-7

GENERAL INFORMATI

APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Botts, John T.
APPLICANT: Juppner, Harald

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PATENT NO. 5460978
; PATENT NO. 5460978
; PATENT MARTIN, THOMAS J.; MOSELEY, JANE M.; KEMP, BRUCE E.; WETTENHALL, RICHARD E.H.
TITLE OF INVENTION: PROTEIN ACTIVE IN HUMORAL ; HYDERCALCEMIA OF MALIGNANCY-PTHRP
NUMBER OF SEQUENCES: 4
; CURRENT APPLICATION DATA;
APPLICATION NUMBER: US/07/715,280
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid TITLE OF INVENTION: Hormone-Related Peptide (PTHrP) FILE REFERENCE: 0609,4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US. 60/105,530
EARLIER APPLICATION NUMBER: US. 60/105,530
EARLIER FILING DATE: 1998-10-22
UNDBER OF SEQ ID NOS: 13
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 7
LENGTH: 9
TABLE OF TABLE OF THE OFFICE OF THE OFFICE OFFI CONTROL OF THE OFFI CO
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Patent No. 6495662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Description of Artificial Sequence: synthetic OTHER INFORMATION: peptide
                                                                                                 Score 35; DB Pred. No. 2.2; 0; Mismatches
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Pred. No. 3e+05;
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Pred. No. 3e+05;
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                                                                                                                                                              DB 6;
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                                                                                                     Indels
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APPLICANT: Kronenberg, Henry
APPLICANT: Fotts, John T.
APPLICANT: Potts, John T.
APPLICANT: Potts, John T.
APPLICANT: Potts, John T.
APPLICANT: Of INVENTION: Bioactive Peptides and Peptide Derivatives of TITLE OF INVENTION: Hormone-Related Peptide (PTHP) and Parathyroid TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570001
CURRENT APPLICATION UNUBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
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LENGTH: 9
TYPE: PRT
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Patent No. 6030790
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/
EILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 0782
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
COMPUTER: PM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: PatentIn Release #1.0, 1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Peptides from the hPTH Sequence NUMBER OF SEQUENCES: 36 CORRESPONDENCE ADDRESS: ADDRESSEE: Jones & Askew, LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: Georgia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Atlanta
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7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             E: Jones & Askew, LLP
191 Peachtree Street,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  USA
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77.8%;
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                                                                                                                                                                                                PCT/EP95/03757
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                                        07826-0007
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Pred. No.
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Query Match
Best Local Similarity
Matches 8; Conserve

Conservative

59.3%;

AVSEIQLMHN 10

5460978-3

SEQ ID NO:3:

LENGTH: 11 FILING DATE: APPLICATION NUMBER: US/ FILING DATE: 14-JUN-1991 PRIOR APPLICATION DATA:

APPLICATION NUMBER: 199,235 FILING DATE: 09-MAY-1988 APPLICATION NUMBER:

RESULT 6 5460978-3

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AVSEIQLLH 9 AVSEIQLMH 9 Conservative

Query Match Best Local

67.8%;

8 Similarity FEATURE:

TYPE: PRT

ORGANISM: Artificial Sequence

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RESULT 10
US-08-817-547A-4
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                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: synthetic;
OTHER INFORMATION: peptide
US-09-421-379-10
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                                                                                                                                                Sequence 4, Application US/08817547A Patent No. 6030790 GENERAL INFORMATION:
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Best Local (
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Best Local (
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APPLICANT: Potts, John T.
APPLICANT: Dotts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bloactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 6069.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US. 60/105,530
EARLIER FILING DATE: 1998-10-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-421-379-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            atent No.
                               APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Mar
TITLE OF INVENTION: Pepti
NUMBER OF SEQUENCES: 36
CORRESPONDENCE ADRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Ver.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Gardella, Thomas J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
HYPOTHETICAL:
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STRANDEDNESS: unl
   ADDRESSEE:
STREET: 19
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                                                                                                                                                                                                                                                                                  1 SVSEHQLLH 9
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                                                                                                                                                                                                                                                                                                                   AVSEIQLMH 9
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3: Jones & Askew, LLP
191 Peachtree Street, 37th Floor
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87.5%;
                                                                          Peptides
                                                                                              Markus
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1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                        Score 30; DB 4;
Pred. No. 3e+05;
                                                                            from
                                                                          the hPTH Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                3e+05;
                                                                                                                                                                                                                                                                                                                                                                                            DB 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 3;
                                                                                                                                                                                                                                                                                                                                                                                          Length 9;
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                                                                                                                                                                                                                                                                                                                                                    Indels
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COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN PC-DOS/MS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/428,257A
FILING DATE: 07/05/95
CLASSIFICATION: 514
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-428-257A-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 14, Application US/08428257A
Patent No. 5885808
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/03757
APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 07826-0007
                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Spooner, Robert A.
APPLICANT: Spenetos, A.A.
TITLE OF INVENTION: Compounds to
NUMBER OF SEQUENCES: 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 7 amino acids
                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Jules E. Goldberg
STREET: 261 Madison Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION: TELEPHONE: 404-818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                STREET: 261 Mac
CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 7 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 10016-2391
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85.7%;
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Pred. No. 3e+0
1; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 3; Length 7; 3e+05;
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                                                                                                                                          (EPO)
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; TOPOLOGY: 1: ; MOLECULE TYPE: US-08-428-257A-14

protein

TYPE: amino acid TOPOLOGY: linear

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RESULT 12
US-08-802-981-124
                                                                                                            RESULT 13
US-08-177-109A-56
                                                                                                                                                                                                                                                                                                                                      ; LOCATION: 3
; OTHER INFORMATION: /product= "Aib" US-08-802-981-124
             Sequence 56, Application US/08177109A
Patent No. 5869615
GENERAL INFORMATION:
APPLICANT: Dennis E. Hourcade and
TITLE OF INVENTION: MODIFIED COMP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/802,981
FILING DATE: 20-FEBB-1997
CLASSIFICATION: 435
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Patent No. 6037137
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                              Best Local Similarity
                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Hunter, Tom
REGISTRATION NUMBER: 38,498
REFERENCE/DOCKET NUMBER: 0166
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NERALL INFORMATION KOMORIYA, Akira
APPLICANT: Komoriya, Akira
APPLICANT: Komoriya, Akira
APPLICANT: Packard, Beverly S.
TITLE OF INVENTION: Compositions for the Detection of Enzyme
TITLE OF INVENTION: Activity in Biological Samples and Methods of Use Thereof
TITLE OF SECUENCES: 231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 231
CORRESPONDENCE ADDRESSE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 11 amino acids
TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Two Embarca
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: Modified-site
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6037137
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                                                                                                                                                                                     QLLHN 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     California
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                                                                                                                                                                                                                                                                              42.4%;
80.0%;
. Hourcade and Teresa J. Oglesby MODIFIED COMPLEMENT PROTEASES 62
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Pred. No. 1.4e+02;
1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      016865-000300US
                                                                                                                                                                                                                                                          Score 25; DB 3; Le
Pred. No. 1.6e+02;
                                                                                                                                                                                                                                                                                                Length 11;
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TOPOLOGY: lir

; MOLECULE TYPE:

; HYPOTHETICAL: N

US-08-177-109A-56
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Best Local Similarity
Whiches 3, Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No. 5928892
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/177,109A
FILING DATE: 03-CAN-1994
CLASSIFETCATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: PABSE, PATENE, 31,284
REFERENCE/DOCKET NUMBER: WU 107
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
ZIP: 30303-3450
ZIP: 30303-3450
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN PC-DOS/MS-DOS
SOFTWARE: PATENTIN PATENTIN PATENT APPLICATION DATA:
APPLICATION NUMBER: US/08/687,706
FILING DATE: 26-JUL-1996
CLASSIFICATION 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/177,109
APPLICATION NUMBER: US 08/177,109
FILING DATE: 03-JAN-1994
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No. 5928892
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (404) 873-87 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
Patrea L. Pabst
STREET: 2800 One Atlantic Center
STREET: 1201 West Peachtree Street
                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Dennis E. TITLE OF INVENTION: NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Patrea L. Pabst
STREET: 2800 One Atlantic Center
STREET: 1201 West Peachtree Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (404) F
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ZIP: 30309-3450
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                                                                                                                                                                                                                                                                                CITY: Atlanta
                                                                                                                                                                                                                                               COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 8 MHNLG 12
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1: USA
                                                                                                                                                                                                                                                               Georgia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Application US/08687706
                                                                                                                                                                                                                                               USA
                                                                                                                                                                                                                                                                                                                                                                                                               Dennis E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          v: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (404) 873-8794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          873-8795
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                                                                                                                                                                                                                                                                                                                                                                                              MODIFIED COMPLEMENT F
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Pred. No. 3e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                PROTEASES
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CLASSIFICATION: 514 ATTORNEY/AGENT INFORMATION:

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LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: peptide
HYPOTHETICAL: no
ANTI-SENS: no
US-08-817-547A-17
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:

MEDDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIAN PC-DOS/MS-DOS
SOFTWARE: PATENTIAN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/817,547A
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 07826-0007
FREERENCE/DOCKET NUMBER: 07826-0007
TELEPHONE: 404-818-3700
TELEPAX: 5 anino acids
TYPE: amino acids
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INFORMATION FOR SEQ ID NO: 56:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 17, Application US/08817547A
Patent No. 6030790
GENERAL INFORMATION:
APPLICANT: Adermann, Knut
APPLICANT: Hock, Dieter
APPLICANT: Hock, Diete
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 40.7%;
Best Local Similarity 60.0%;
Matches 3; Conservative
Query Match 40.7%; Score 24; DB 3; Best Local Similarity 100.0%; Pred. No. 3e+05; Matches 4; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 31,284
REFERENCE/DOCKET NUMBER: WU 107 DIV
TELECOMMUNICATION INFORMATION:
TELEPHONE: (404) 873-8794
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 191 Peac
CITY: Atlanta
STATE: Georgia
COUNTRY: USA
ZIP: 30303
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:||:|
1 LHNMG 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          E: Jones & Askew, LLP
191 Peachtree Street, 37th Floor
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Pred. No. 3e+05;
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Qy 9 HNLG 12 |||| Db 1 HNLG 4

Search completed: May 18, 2004, 10:06:44 Job time: 12.75 secs

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seq length: 12
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T-cell receptor al	potB protein - Sal	transcription fact	beta-D-galactosida	retinal oxidase -	protein QA300042 -	endo-1,4-beta-xyla	translation elonga	calliFMRFamide 12	e antigen p20e pre	ranatachykinin C -		neuromedin K - pig	1	chlorophyll a/b-b1	neuropeptide Led-C

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cytochrome b559 component psbF - pepper chloroplast (fragment)
C;Species: chloroplast Capsicum annuum (pepper)
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 11-Jun-1999
C;Accession: S28055
R;Kuntz, M.; Camara, B.; Weil, J.H.; Schantz, R.
Plant Mol. Biol. 20, 1185-1188, 1992
A;Title: The psbt gene from bell pepper (Capsicum annuum): plastid RNA editing A;Reference number: S28055; MUID:93099270; PMID:1463853
A;Accession: S28055; MUID:93099270; PMID:1463853
                                                                                                                                                                                                                                       RESULT 2
PHILE7
T-cell receptor alpha chain V region (Cw3/IFI1) - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 16-Jul-1999
Sequence_revision 16-Jul-1999
Htext_change 16-Jul-1999
C;Accession: PHILE7
R;Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, A.; Gournier, H.; Barra, C.; Wic
J. Exp. Med. 176, 439-447, 1992
A;Title: H-2-restricted cytolytic T lymphocytes specific for HLA display T cell receptor
A;Reference number: $26512; MUID:92364546; PMID:1380061
A;Accession: PHILE7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Genome: chloroplast
C;Superfamily: cytochrome b559 component
C;Keywords: chloroplast; photosynthesis;
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A;Residues: 1-10 <KUN>
A;Cross-references: EMBL:X65570; NID:g14344; PIDN:CAA46539.1; PID:g415734
C;Genetics:
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A;Molecule type: mRNA
A;Residues: 1-12 <CAS>
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37.5%; Pred. No. 7.4e+0:
tive 4; Mismatches
                                                                                                           32.8%;
                                                                                      Score 20; DB 2;
Pred. No. 9.1e+02;
0; Mismatches 2
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A; Residues:
C; Kevw~~
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A;Accission: $78420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Stoffels, G.; Nissen-Meyer, J.; Gudmundsdottir, A.; Sletten, K.; Holo, H.; Nes, I.F. Appl. Environ. Microbiol. 58, 1417-1422, 1992
A;Title: Purification and characterization of a new bacteriocin isolated from a Carnobac A;Reference number: A58718; MUID:92321768; PMID:1622206
A;Accession: A58718
                                                                                                                                                                                                                           C;Accession: A60589
R;Yoshino, K.I.; Kajiura, H.; Nomura, K.; Takao, T.; Shimonishi, Y.; Kurita, M.; Yamagud Comp. Biochem. Physiol. B 94, 739-751, 1989
A;Title: A halogenated amino acid-containing sperm activating peptide and its related peotus nudus, Echinometra mathaei and Heterocentrotus mammillatus.
₽.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        C;Species: Rattus norvegicus (Norway rat)
C;Date: 25-Peb-1998 #sequence revision 13-Mar-1998
C;Accession: S78420
R;Goldschmidt-Reisin, S.; Graack, H.R.
                             Š
                                                                                                                                                                                                                                                                                                                        sperm-activating peptide (Tyr-2, Asn-3, Gly-5, Ile-9, Asp-10 SAP-I) - slate-pencil urchi
C;Species: Heterocentrotus mamillatus
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 18-Aug-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             C;Species: Carnobacterium sp.
C;Date: 23-Jan-1998 #seguence_revision 30-Jan-1998 #text_change
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A;Residues: 1-10 <YOS>
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Pred. No. 2.8e+05;
0; Mismatches 0;
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Pred. No. 1.8e+03;
1; Mismatches 0
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Pred. No. 2.8e+05;
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T-cell receptor alpha chain V region (CW3/10.1) - human (fragment) C;Species: Homo sapiens (man) C;Species: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16:C;Accession: PH1190 #sequence_revision 16-Jul-1999 #text_change 16:C;Accession: PH1190 #sequence_revision 16-Jul-1999 #text_change 16:C;Accession: PH1190 #sequence_revision 16-Jul-1999 #text_change 16:Display #sequence_revision 16:Display 
                                                                                                                                                                                                                                                                                                                                                                                                                              R,Ruff-Jamison, S.; Campos-Gonzalez, R.; Glenney Jr., J.R.
J. Biol. Chem. 266, 6607-6613, 1991
A;Title: Heavy and light chain variable region sequences and antibody properties of anti-A;Reference number: A38740; MUID:91177923; PMID:1706720
A;Reference number: A38740; MUID:91177923; PMID:1706720
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A;Title: Structures of two cockroach neuropeptides assigned by fast atom bombardment A;Reference number: A90118; MUID:85046530; PMID:6548628
A;Recession: A05169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  neuropeptide M-I - American cockroach (Species: Periplaneta americana (American cockroach) (Date: 05-Jun-1987 #sequence_revision 05-Jun-1987 #text_change 31-Dec-1993 (Accession: A05169 R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; O'Shea, M.; Cook, J.C.; Hemling, M.E.; Rine: R;Witten, J.L.; Schaffer, M.H.; Rine: R;Witten, M. R;Wi
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A05169
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A37268
C;Accession: A37268
C;C;Accession: A37268
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A; Residues: 1-12 < CAS>
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C; Keywords: neuropeptide
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                                                                                                                                                                                                                                                                                                                                        A;Status: preliminary
A;Molecule type: mRNA
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Pred. No. 2.8e+05;
1; Mismatches 3
                                                                                           Score 17; DB 2;
Pred. No. 2.9e+03;
1; Mismatches 4
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Bur. J. Blochem. 205, 341-345, 1992
A;Title: Placental alkaline phosphatase has a binding site A;Reference number: S21205; MUID:92209522; PMID:1555592
A;Accession: S21205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ig heavy chain V region - human
C;Species: Homo sapiens (man)
C;Date: 22-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 16-Aug-1996
C;Accession: S21205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Molecule type: protein
A;Residues: 1-12 <SCH>
A;Residues: 1-12 <SCH>
A;Rote: the amino end of this peptide is not blocked
A;Note: synthetic locustamyotropin mimics natural locustamyotropin only in the C;Comment: This peptide was shown to stimulate visceral muscle contractions in C;Superfamily: pyrokinin
C;Keywords: amidated carboxyl end; neuropeptide
F;12/Modified site: amidated carboxyl end (Leu) #status experimental
                                                              A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-12 <CAS>
                                                                                                                          C;Accession: PH1189

R;Casanova, J.L.; Cerottini, J.C.; Matthes, M.; Necker, P. J. Exp. Med. 176, 439-447, 1992

A;Title: H-2-restricted cytolytic T lymphocytes specific A;Reference number: S26512; MUID:92364546; PMID:1380061

A;Accession: PH1189
                                                                                                                                                                                                                                                      T-cell receptor alpha chain V region (Cw3/2C3) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
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A;Molecule type: protein
A;Residues: 1-12 <MAK>
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R;Schoofs, L.; Holman, G.M.; Hayes, T.K.; Tips, A.; Nachman, R.J.; Va Peptides 11, 427-433, 1990
A;Title: Isolation, identification and synthesis of locustamyotropin A;Reference number: A43975; MUID:90341077; PMID:1974346
A;Accession: A43975
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     N;Alternate names: Lom-MT
C;Species: Locusta migratoria (migratory locust)
C;Date: 11-Feb-1993 #sequence_revision 02-Jun-1994 #text_change 08-Dec-1995
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Pred. No. 3.5e+03;
3; Mismatches 3;
Score 17;
Pred. No.
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DB 2; L
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RESULT S09138

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A;Molecule type: protein
A;Residues: 1-10 <GASI>
R;Gaede, G.; Rinehart Jr., K.L.
Biol. Chem. Hoppe-Seyler 368, 67-75, 1987
Biol. Chem. Hoppe-Seyler of the hypertrehalosaemic factor II from the A;Title: Primary structure of the hypertrehalosaemic factor II from the A;Reference number: S07157; MUID:87157103; PMID:3828078
A;Accession: S07157
A;Molecule type: protein
A;Residues: 'Z',2-10 <GABZ>
A;Residues: 'Z',2-10 <GABZ>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Carausius morosus
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: JC11416; 807157
R;Gaede, G; Kellner, R.; Rinehart, K.L.; Proefke, M.L.
Biochen. Biophys. Res. Commun. 189, 1303-1309, 1992
A;Title: A tryptophan-substituted member of the AKH/RPCH family isolated from a stick in A;Reference number: JC1416; MUID:93129188; PMID:1482345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R;Friedman, D.J.; Umeda, P.K.; Sinha, A.M.; Hsu, H. Proc. Natl. Acad. Sci. U.S.A. 81, 3044-3048, 1994 A;Title: Characterization of genomic clones specifying A;Reference number: I46888; MUID:84221901; PMID:8328491 A;Accession: I46868
                                                                                                                           F;8/Binding site: carboxylend (Thr) F;10/Modified site: amidated carboxyl end (Thr)
                                                                                                                                                                    C;Keywords: amidated carboxyl end; glycoprotein; hormone; neuropeptide; p F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental F;8/Binding site: carbohydrate (Trp) (covalent) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hypertrehalosemic hormone I - stick insect N;Alternate names: neuropeptide Cam-HrTH-I N;Contains: hypertrehalosemic factor II
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      alpha-myosin heavy chain - rabbit (fragment)
C;Species Cryyttolagus cuniculus (domestic rabbit)
C;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change 05-Nov-1999
C;Accession: 146868
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT
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A; Residues: 1-7 <FRI>
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                                                                       Best
                                                                                                                                                                                                                                            Comment: Hypertrehalosemic factor II lacks the tryptophan modification.
Comment: This peptide raises hemolymph levels of trehalose in the cockroach Periplaneus Superfamily: adipokinetic hormone
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Pred. No. 2.8e+05;
1; Mismatches 1
                                                   Score 16; DB
Pred. No. 4.5e
2; Mismatches
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amine oxidase (copper-containing) (EC 1.4.3.6) - pig (fragment) C;Species: Sus scrofa domestica (domestic pig) C;Date: 12-Oct-1989 #sequence_revision 31-Dec-1993 #text_change 06-Sep-1996 C;Accession: A32428 R;van der Meer, R.A.; van Wassenaar, P.D.; van Brouwershaven, J.H.; Duine, J.A. Biochem. Biophys. Res. Commun. 159, 726-733, 1989 A;Title: Primary structure of a pyrroloquinoline quinone (PQQ) containing pepti A;Reference number: A32428; MUID:89193662; PMID:2539124 A;Accession: A32428
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 15
A32428
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R;Gaede, G.; Rinehart, K.L.
Biol. Chem. Hoppe-Seyler 371, 345-354, 1990
Biol. Chem. Hoppe-Seyler 371, 345-354, 1990
A;Title: Primary structures of hypertrehalosaemic neuropeptides isolated from the corpor entalis and of the stick insect Extatosoma tiaratum assigned by tandem fast atom bombard A;Reference number: S08995; MUID:90253659; PMID:2340112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            hypertrehalosemic hormone II - stick insect (Extatosoma tiaratum)
N;Alternate names: Cam-HrTH-II
C;Species: Extatosoma tiaratum
C;Species: 30-Jun-1992 #sequence_revision 24-Oct-1997 #text_change 31-Oct-1997
Search completed: May 18, 2004, 10:05:41 Job time: 10.25 secs
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A; Residues: 1-7, 'K', 9-11 < VAN>
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Copyright (c) 1993 - 2004 Compugen Ltd.
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P36960 carnobacter
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10-OCT-2003
RO I (Hypert
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MOD_RES 1 1 PYRE
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SEQUENCE 10 AA; 1163 MW; 0562
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE.
TISSUE-Corpora cardiaca;
TISSUE-Corpora cardiaca;
TISSUE-Side of two neuropeptides of the AKH/RPCH-family
"Sequence analyses of two neuropeptides of the AKH/RPCH-family
the lubber grasshopper, Romalea microptera.";
Peptides 9,681-688(1988)
-!- FUNCTION: Hypertrehalosaemic factors are neuropeptides that
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            a Carnobacterium sp.";
Appl. Environ. Microbiol. 58:1417-1422(1992)
-!- FUNCTION: Lanthionine-containing peptide
Active on Gram-positive bacteria.
Antibiotic; Bacteriocin; Lantibiotic.
                                                                                                                                                                                                                                                                                                                                         + +
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Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridomorpha;
Acridoidea; Romaleidae; Romalea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=92321768; PubMed=1622206; Stoffels G., Nissen-Meyer J., Gudmundsdottir
                                                                                                                                                                                                                                                                    InterPro; IPR002047; AKH.
PROSITE; PS00256; AKH; 1.
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Ol-JUN-1994 (Rel. 29, Last sequence uj
16-OCT-2001 (Rel. 40, Last annotation
Lantibiotic carnocin UI49 (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P36960;
01-JUN-1994
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                                                                                                                                                                                                                                                                                                                          FUNCTION: Hypertrehalosaemic factors are neuropeptides that elevate the level of trehalose in the hemolymph (trehalose is major carbohydrate in the hemolymph of insotym). SUBCELLULAR LOCATION: Secreted.
SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
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BB-1994 (Rel. 28, Last sequence up
CT-2003 (Rel. 42, Last annotation
(Hypertrehalosaemic factor).
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   Conservative
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                                                                                       Duve H., Johnsen A.H., Maest
Davey M., East P.D., Thorpe
"Lepidopteran peptides of th
Peptides 18:1301-1309(1997)
-!- SIMILARITY: Belongs to
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Neuropeptide; Amidation.
MOD_RES 8 8
SEQUENCE 8 AA; 936 MW
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MOD_RES 12 12 AMIDA
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Neoptera; Orthopteroidea; Orthoptera; Caelifera; Acridomorpha;
Acridoidea; Acrididae; Oedipodinae; Locusta.
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01-AUG-1991 (Rel. 19, Last annotation
Locustamyotropin 1 (LOM-MT-1).
Locusta migratoria (Migratory locust)
                                                                                                                                                                                                                        MEDLINE=98054539;
Duve H., Johnsen F
                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cydia pomonella (Codling moth)
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PROSITE; PS00539; PYROKININ; 1.
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TISSUE=Corpora cardiaca;
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                                                                                                                                                                                                                                                                                                 TISSUE=Larva;
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      936 MW;
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Pred. No. 2.
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                                                                                                                                                            superfamily."
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RESULT 6
HTF2 CARMO
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Best Local
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Best Local (
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MEDLINE-33129188; PubMed=1482345;
Gaede G., Kellner R., Rinehart K.L. Jr., Proefke M.L.;
"A tryptophan-substituted member of the AKH/RPCH family isolated a stick insect corpus cardiacum.";
Biochem. Biophys. Res. Commun. 189:1303-1309(1992).
-!- FUNCTION: Hypertrehalosaemic factors are neuropeptides that elevate the level of trehalose in the hemolymph (trehalose i major carbohydrate in the hemolymph of insects).
-!- SUBCELTULAR LOCATION: Secreted.
-!- MASS SPECTROMETRY: MW=1308.61; METHOD=FAB.
-!- SIMILARITY: Belongs to the AKH / HRTH / RPCH family.
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                                                                            CARBOHYD
MOD_RES
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                      the corpora cardiaca of the cockroaches Leucophaea maderae, Gromphadorhina portentosa, Blattella germanica and Blatta orientalis and of the stick insect Extatosoma tiaratum assigned by tandem fast atom bombardment mass spectrometry."; Blook. Chem. Hoppe-Seyler 371:345-354(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                       Gaede G., Rinehart K.L. Jr.;
"Primary structures of hypertrehalosaemic neuropeptides isolated from the corpora cardiaca of the cockroaches Leucophaea maderae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Primary structure of the hypertrehalosaemic factor II from the corpus cardiacum of the Indian stick insect, Carausius morosus, determined by fast atom bombardment mass spectrometry."; Biol. Chem. Hoppe-Seyler 368:67-75(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SPECIES=C.morosus; TISSUE=Corpora cardiaca; MEDLINE=87157103; PubMed=3828078; Gaede G., Rinehart K.L. Jr.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Extatosoma tiaratum (Stick insect).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroidea; Phasmatodea; Euphasmida;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-JUI-1989 (Rel. 11, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
10-CCT-2003 (Rel. 42, Last annotation updat
Hypertrehalosaemic factor II (HTF-II) (HRTH
                                                                                       Neuropeptide; Amidation; Glycoprotein; Pyrrolidone carboxylic MOD RES 1 PYRROLLDONE CARBOXYLIC ACID. CARBOHYD 8 8 C-LINED (MAN) (PROBABLE). MOD_RES 10 10 AMIDATION.
                                                                                                                                                                   PIR;
                                                                                                                                                                                                                                                                                                                              CARBOHYDRATE-LINKAGE SITE.
CDECTES=C.morosus; TISSUE=Corpora cardiaca;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              SPECIES=E.tiaratum; TISSUE=Corpora MEDLINE=90253659; PubMed=2340112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                neuropeptide II).
Carausius morosus (Indian stick insect),
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S09138; S09138.
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  EIQFMHNLG
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                                                                            1164 MW;
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Pred. No.
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Pred. No. 2.9e
2; Mismatches
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Phasmatoidea;
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RESULT 9
EFG_CLOPA
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RESULT 7
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Submitted (JUL-1997) to Swiss-Prot
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P81010;
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15-JUL-1998 (Rel. 36, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation updat
Allergen Fus s 13596* (Fragment)
Fusarium solani (subsp. pisi) (Nectria haem
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Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea;
Pelodryadinae; Litoria.
Pelodryadinae; Litoria.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Electrin
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10-OCT-2003
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Hypocreomycetidae; Hypocreales; Nectriaceae; Nectria
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MOD_RES
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TISSUE SPECIFICITY: Skin.
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(Rel. 42, Last annotation update)
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allergic re
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668761F2C9A00000 CRC64;
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Pred. No. 1.4e
1; Mismatches
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Pred. No.
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                                                                                 Mismatches
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Australian buzzing tree frog
skin peptides from Litoria
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1.4e+05;
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Hylidae;
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RESULT 10
NUHM_CANFA
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              Query Match
Best Local
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P81350;
15-JUL-1998
15-JUL-1998
28-FEB-2003
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"Two-dimensional gel electrophoresis separation and N-
sequence analysis of proteins from Clostridium pasteur
Electrophoresis 19:802-806(1998).

-i- FUNCTION: This protein promotes the GTP-dependent
                                                                                                                                                                                                                                                      NUHM CANFA
P49820;
                                                                                                                                                                                                                                                                                                                                                                                      PROSITE; PS00301; EFACTOR GTP; PARTIAL.
Elongation factor; Protein biosynthesis;
NON_TER 11
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Bacteria; Firmicutes; Clo
                                                                                                                                                                                                                   01-OCT-1996 (Rel. 34, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
10-CCT-2003 (Rel. 42, Last annotation update)
NADH-ubiquinone_oxidoreductase 24 kDa subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Elongation
                                                                                                                                                 SEQUENCE
                                                                                                                                                                       Canis familiaris (Dog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Carnivora; Fissipedia; Canidae;
                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE
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                                                                                                                                        TISSUE=Heart;
                                                                                                                                                                                                                                                                                                                                                                                                                SUBCELLULAR LOCATION: Cytoplasmic SIMILARITY: Belongs to the GTP-bit EF-G/EP-2 subfamily.
erPro; IPR000795; EF-GTPbind.
SIMILARITY:
                                                                                                                                                                                                            1.6.99.3) (Fragment).
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          mitochondrial
                                                                                                                                                                 _TaxID=9615;
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h factor
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Belongs to the
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          inner membrane.
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Pred. No. 5.1e
3; Mismatches
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complex I
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pasteurianum W5
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Canis.
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RESULT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE, AND MASS SPECTROMETRY.

MEDILINE=90319122; PubMed=1973541;

Grimmelikhuijzen C.J.P., Rinehart K.L. Jr., Jacob E., Graff D.,

Reinscheid R.K., Nothacker H.-P., Staley A.L.;

"Isolation of L-3-phenyllactyl-Leu-Arg-Asn-NH2 (Antho-RNamide),
anemone neuropeptide containing an unusual amino-terminal block:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anthopleura elegantissima (Sea anemone)
Eukaryota; Metazoa; Cnidaria; Anthozoa;
Nynantheae; Actiniidae; Anthopleura.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-FEB-2003 (Rel. 41,
28-FEB-2003 (Rel. 41,
10-OCT-2003 (Rel. 42,
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P58707;
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NON_TER 11
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Proc. Natl.
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                                                                                                                                                    Eukaryota; Metazoa; Arthropoda; Hexapoda; Insec
Neoptera; Endopterygota; Lepidoptera; Glossata;
Tortricoidea; Tortricidae; Olethreutinae; Cydia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Neuropeptide; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NCBI_TaxID=6110;
rissue=Larva;
MEDLINE=98054539;
                                                                    SEQUENCE
                                                                                                                                                                                                                                           Cydia pomonella (Codling
                                                                                                                                                                                                                                                                       Cydiastatin
                                                                                                                        NCBI_TaxID=82600;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MASS SPECTROMETRY: Neuron specific.
MASS SPECTROMETRY: NW=549.3; METHOD=FAB
A35779; A35779.
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AMIDATION.
64540729A0000000 CRC64;
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1; Mismatches
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Pred. No. 1.4e+05;
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.on update)
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PubMed=9392829

SO CCC RTT RANGE

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RESULT 13
TYS1_LITRU STANDARD; PRT; 7 AA.
AC P82065;
DT 28-FEBS-2003 (Rel. 41, Last sequence update)
DT 10-CTT-2003 (Rel. 42, Last annotation update)
DT 10-CTT-2003 (Rel. 42, Last annotation update)
DT Tryptophyllin 5.1.
OS Litoria rubella (Desert tree frog).
Litoria rubella (Desert tree frog).
CC palodryddinae; Litoria.
OX NCBI_TaxID=104895;
RN [1]
RP SEQUENCE, AND MASS SPECTROMETRY.
RC TISSUE-Skin secretion;
RC TISSUE-Skin secretion;
RA Tyler M.J., Wallace J.C.;
RA Tyler M.J., Wallace J.C.;
RA Tyler M.J. Wallace J.C.;
RI (Litoria rubella'. The skin peptide profile as of evolutionary trends of amphibians.";
RL Aust. J. Chem 49:955-963(1996).
CC -!- FUNCTION: May act as a neuromodulator or no CC -!- FUNCTION: May act as a neuromodulator or no CC -!- FUNCTION: May act as a neuromodulator or no CC -!- FUNCTION: May act as a neuromodulator or no CC -!- FUNCTION: May act as a neuromodulator or no CC -!- FUNCTION: May act as a neuromodulator or no CC -!- FUNCTION: May act as a neuromodulator or no CC -!- FUNCTION: May act as a neuromodulator or no CC -!- FUNCTION: May act as a neuromodulator or no CC -!- FUNCTION: May act as a neuromodulator or no CC -!- FUNCTION: Secreted.
CC -!- FUNCTION: May act as a neuromodulator or no CC -!- FUNCTION: Secreted.
CC -!- MASS SPECTROMETRY: Mw=965; METHOD-FAB.
FM MOD_RES 7 AA; 983 MM; 7401E9D3676046BO CR
RESULT 14
AL18 CARMA
ID AL18 C
AC P81821
DT 30-MAY
DT 30-MAY
DT CARCIN
OS CARCIN
OS CARCIN
OC EUMA12
OC EUMA12
OC EUMA16
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P81821;
30-MAY-2000
30-MAY-2000
30-MAY-2000
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Litoria rubella (Desert tree frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Amphibia; Batrachia; Anura; Neobatrachia; Hyloidea;
pelodryadinae; Litoria.
NCBI TaxID=104895;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Winstan. Davey M., East P.D., Thorpe A., "Lepidopteran peptides of the allatostatin superfamily.", Peptides 18:1301-1309(1997).

-!- SIMILARITY: Belongs to the allatostatin family.
                                            Carcinustatin 18.

Carcinus maenas (Common shore crab) (Green crab).

Custacea; Malacostraca;

Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;

Eukaryota; Metazoa; Bucarida; Decapoda; Pleocyemata; Brachyu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Neuropeptide; Amidation. MOD_RES 7 7
                  Eukaryota; Metazoa; Arthropoda;
Eumalacostraca; Eucarida; Decapo
Eubrachyura; Portunoidea; Portun
NCBI_TaxID=6759;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Amphibian defense peptide; Amidation; Neuropeptide; Pyrrolidone carboxylic acid.
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Best Local
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Davey M., East P.D., Thorpe A.;
"Lepidopteran peptides of the allatostatin sup
Peptides 18:1301-1309(1997).
- SIMILARITY: Belongs to the allatostatin fa
Neuropeptide; Amidation.

NOD RES

8

AMIDATION.
                                                                                                                                 Query Match
Best Local
                                                                                                            Matches
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P82152;
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Cydiastatin
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Eur. J. Blochem. 250:727-734(1997).
Eur. FUNCTION: May act as a neurotransmitter
-i- SIMILARITY: Belongs to the allatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cydia pomonella (Codling moth).

Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta; Pterygota;
Neoptera; Endopterygota; Lepidoptera; Glossata; Ditrysia;
Tortricoidea; Tortricidae; Olethreutinae; Cydia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE.
TISSUE=Cerebral ganglion, and Thoracic ganglion;
MEDLINE=98121193; PubMed=9461295;
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MOD_RES 8 AMIDATION (POTENTIAL)
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                                                                                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rissum=Larva;
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(Rel. 39, Last sequence update)
(Rel. 39, Last annotation update)
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L; Mismatches 1
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 12
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Perfect score:
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                                                                                                                                                                                                                                                  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                            Score
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125.164 Million cell updates/sec
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Match
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1: sp_archea:*
2: sp_bacteria:*
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Gapop 10.0 , Gapext 0.5
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 GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd.
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sp_phage:*
sp_plant:*
sp_virus:*
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sp_unclassified:*
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Q9TT78
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Q82085 limnodynast
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ALIGNMENTS

PRELIMINARY;

PRT;

11 AA

RESULT QSTT78 ID Q AC Q DT Q	Qy da	Query Best Match				2888	
LT 2 Q9TT78 PRELIMINARY; PRT; 8 AA. Q9TT76; Q9TT76; 01-MAY-2000 (TrEMBLrel. 13, Created) 01-MAY-2000 (TrEMBLrel. 13, Last sequence update)	9 HNLG 12 7 HNLG 10	Query Match 39.3%; Score 24; DB 13; Length 11; Best Local Similarity 100.0%; Pred. No. 6.3e+02; Matches 4; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	Arch. Biochem. Biophys. 289:197-207(1991). NON TER 11 11 SEQUENCE 11 AA; 1209 MW; 7CA02D1D41E8772B CRC64;	Guan A.L., Retzios A.D., Henderson G.N., Markland F.S.dr.; "Purification and characterization of a fibrinolytic enzyme from venom of the southern copperhead snake (Agkistrodon contortrix contortrix).";	[1] SEQUENCE. MEDLINE=91378546; PubMed=1898066;	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea; Viperidae; Crotalinae; Agkistrodon. NCBT TaxTD=8720.	ce update) tion update) t).

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Eukaryota; Metazoa; Chordata; C
Mammalia; Eutheria; Carnivora;
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Viruses; ssRNA negative-strand viruses; Mononegavirales;
Paramyxoviridae; Paramyxovirinae; Morbillivirus.
NCBI_TaxID=11232;
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Brouillette J.A.,
Canine distemper virus.
Viruses; ssRNA negative-strand
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Liermann H., Harder T., Haas
"Genetic analysis of the cen
proximal coding part of the
morbilliviruses.";
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Brouillette J.A., Andrew J.R., Venta
"Estimate of nucleotide diversity in
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O13591;
O1-JAN-1998 (TrEMBLrel. 05, C
O1-NOV-1999 (TrEMBLrel. 12, I
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ORF YNL337W (Fragment).
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STRAIN=Dog #10757/96;

Grenann H., Harder T., Haas L.;

Genetic analysis of the central untranslated genome region and "Genetic analysis of the F gene of wild-type and vaccine dist morbillivituses.";

Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
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RAFLERY M.J., Bradford A.M., Bow:
Reptides from Australian frogs.
"Peptides from Simmodynastes in
the banjo frogs Limmodynastes in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Limnodynastes terraereginae (Northen banjo frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Myobatrachi
Limnodynastinae; Limnodynastes.
                                                                                                                                                                                                                                        Saccharomyces cerevisiae (Baker's yeast).

Bukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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                                           SEQUENCE FROM N.A.
Obermaier B., Piravandi
Submitted (MAY-1996) to
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SPECTROMETRY: MW=1236; METHOD=FAB.
12 AA; 1236 MW; 147AA70FDF472724
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noma virus MH2
s Y73 and SR-A.
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01-MAY-2000
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Q23876;
Q1-NOV-1996
Q1-NOV-1996
Q1-NOV-1998
                                                                                                         Bradford A.M., Raftery M.J., Bowie J.H., We Peptides from Australian frogs. The struct Limnodynastes salmini and Fleberin from I Aust. J. Chem. 46:1235-1244(1993).

-!- MASS SPECTROMETRY: MW-1114; METHOD=FAB. SEQUENCE 12 AA; 1114 MW; 3AB5A976CAA727
                                                                                                                                                                                                                                                Limnodynastes salmini (Salmin's-striped frog).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Myobatrachidae;

Limnodynastinae; Limnodynastes.
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Eukaryota; Mycetozoa; Dic
                                                                                                                                                                                                                                   NCBI_TaxID=39404;
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                                                                                                                                                                                         TISSUE-SKIN SECRETION;
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Mycetozoa; Dictyosteliida;
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46:495-505(1982)
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01-NOV-1996
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SEQUENCE
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MEDLINE=84221901; Pubm
Friedman D.J., Umeda P
Frabinowitz M.;
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EMBL; X83121; CAA58183.1;
MON TER 8 8 770 MW.
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation update)
Alpha-myosin heavy chain (Fragment).
Oryctolagus cuniculus (Rabbit).
Dukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-NOV-1996 (TYEMBLEEL 01, Created)
01-NOV-1996 (TYEMBLEEL 01, Last sequence update)
01-OCT-2002 (TYEMBLEEL 22, Last annotation update)
DNA for ORF'S from chromosome XV (Fragment).
Saccharomyces cerevisiae (Baker's yeast).
Eukaryota, Fungi, Ascomycota, Saccharomycotina, Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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Q28742;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Zumstein E., Pearson B.M., Kalogeropoulos A., "A 29.425 kb segment on the left arm of yeast more than twice as many unknown as known open Yeast 11:975-986(1995).
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STRAIN=FY1679;
MEDLINE=95021609; PubMed=8533473;
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Proc. Natl. Acad. Sci. U.S.A
EMBL; K01698; AAA31415.1; -.
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eda P.K., Sinha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               916 MW;
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U.S.A. 81:3044-3048(1984)
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Last sequence update)
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1e+06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 8;
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RESULT 15
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DYNASTIN 5.
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01-MAY-2000
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                                                                                                                                                                                                                                                      P82083;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bradford A.M., Raftery M.J., Bowie J.H., Wallace J.C., Tyler M.J. Peptides from Australian frogs. The structure of the dynastins Limnodynastes salmini and Fletcherin from Limnodynastes fletcher Aust. J. Chem. 46:1235-1244(1993).

-i- MASS SPECTROMETRY: MW=772; METHOD=FAB.
SEQUENCE 8 AA; 772 MW; 7BS8772455A2C728 CRC64;
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                              Limnodynastes salmini (Salmin's-striped f. Eukaryota; Metazoa, Chordata, Craniata; W. Amphibia, Batrachia, Anura, Neobatrachia,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Limnodynastinae; Limnodynastes. NCBI_TaxID=39404;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Limnodynastes salmini (Salmin's-striped frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Myobatrachidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gramene; Q40659; -.
NON TER 8
SEQUENCE 8 AA; 9:
      Limnodynastinae; Limnodynastes
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Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Alpha-amylase (Fragment).
Oryza sativa (Rice)
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Rodriquez R.L.
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MEDLINE=91078641; PubM
Kumagai M.H., Shah M.,
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Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-OCT-2002 (TrEMBLrel. 22,
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h M., Terashima
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                                     Myobatrachidae;
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OX NCBI_TaxID=39404;

RV [1]

RP SEQUENCE, AND MASS SPECTROMETRY.

RC TISSUE-SKIN SECRETION;

RA Bradford A.M., Raftery M.J., Bowie J.H., Wallace J.C., Tyler M.J.;

RA Bradford A.M., Raftery M.J., Bowie J.H., Wallace J.C., Tyler M.J.;

RT Limnodynastes salmini and Fletcherin from Limnodynastes fletcheri.";

RL Aust. J. Chem. 46:1235-1244(1933).

CC -!- MASS SPECTROMETRY: MW-786; METHOD=FAB.

SQ SEQUENCE 8 AA; 786 MW; 7B58772455B05728 CRC64;

Query Match
Best Local Similarity 100.0%; Pred. No. 1e+06;
Matches 3; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 10 NLG 12

Db SNLG 7

Search completed: May 18, 2004, 10:04:48

Job time: 30.25 secs
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Result
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Maximum
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length: 12
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Match
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                                                                                                                                                                                                                                                                                                                                     A_Geneseq_29Jan04:*
1: geneseqp1980s:*
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geneseqp2003bs:*
geneseqp2004s:*
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Match 100%
first 45 summaries
GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
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         AAR91644

AAV68767

AAB86219

ABAR44166

AAR96225

AAB96225

AAB969666

AAR996966

AAR996966

AAR9969163

AAP9691863

AAP96917

AAB96917

AAB96917

AAB96917

AAB96917

AAB96917
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ABG72608
AAY96968
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                             Aab96931
Aab96931
Aab96915
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Aay96981
Aab86220
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Aab86225
Aab01862
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Aab86219
Abr44166
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Abg72608
Aay96968
Aar91644
                                                                                                   Aay96966
Aar91645
               Abp71484
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                      Parathyro
Human par
PTH(1-14)
Parathyro
Human par
Rat parat
Rat parathyro
Parathyro
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Parathyro
Parathyro
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Parathyro
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(Asn10, T
                                                                                                                  Parathyro
Parathyro
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Human par
Amino aci
Human par
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Resin bou
Human par
PTH(1-14)
parat
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Human; parathyroid hormone; antigen; hypercalcaemia; osteoporosis; primary hyperparathyroidism; mouse; rat; bovine; porcine; canine.

Parathyroid hormone antigenic peptide 2-12.

11-FEB-2003 ABG72607;

(first

entry)

	4. U	44	43	42	41	40	39	38	37	36	G:	34	33	32	31	30	29	28	27	26	
	25	25	25	26	26	27	27	27	27	27	28	28	28	29	29	29	31	31	31	31	
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	Q	σ	σ	w	w	w	N	N	4.	4	4	w	N	w	w	w	D	4.	ð	Q	
AT TOWNSHIP	ABP88771	ABP89380	34	AAB01866		AAY86894	AAY55148	42	ABU54029	AAG63991	AAB86221	O)	AAR91646		AAY78849	AAB01864		AAB96891	ABP71483	ABP71485	-
	p88771 HLA prot	89380 HLA prot	54 HLA		b00068 N-termin	y86894 Human ha	Aay55148 Anti CD34	90421 Antibo	u54029	91 Comple	1 Human pa	07467 Antige	91	7062 F	849 Parathy	01864 PT	482 Parathy	891 Rat	p71483 Par	Abp71485 Parathyro	

ALIGNMENTS

ABG72607 standard; peptide; 11

B

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RESULT 1
ABG72607
ID ABG72607
XX ABG7
AC ABG7
XX Huma
XX Huma
XX Homc
OS Mus
SS Sus
SS Sus
SS Sus
SS Sus
Cani
XX US2(
PN US2(
XX O5-I
XX O5-I
XX ABG7
PR O5-I
XX ABG7
PR O5-I
XX US2(
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Rattus sp.
Bos taurus.
Sus scrofa.
                                                                                                                                                                                                                 New parathyroid hormone (PTH) antigenic peptide inducing the formation and isolation of antibodies having an affinity to it, useful for determining bioactive PTH levels in serum, plasma and/or cell culture
                                                                                                                                                                                                                                                                                                                                                                        WPI;
                                                                                                            Claim 1; Page 5; 11pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                       Zahradnik RJ, Lavigne JR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-DEC-2000; 2000US-00730174.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               05-DEC-2000; 2000US-00730174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ZAHR/) ZAHRADNIK R J.
(LAVI/) LAVIGNE J R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-AUG-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US2002110871-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                        2003-066685/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /label= Leu, Phe
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The invention formation and

relates to a new antigenic peptide for inducing the isolation of antibodies having an affinity to it, being

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RESULT 2
ABG72608
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CC are; (1) a method for producing antibodies useful in the determination of CPTH levels in a biological sample comprising: (a) providing at least one Crists peptide antigen comprising a peptide fragment of PTH; (b) CC administering the first peptide antigen to a host animal to induce CC antibody production; (c) monitoring the antibody titre produced; (d) CC isolating antisera produced in the host animal; and (e) selecting CC antisera from the isolated antisera produced in the host that is capable CC of binding to a second peptide antigen; (2) an antibody (ab) produced by the method; and (3) test kits and analytical procedures used for the CC compositions of the present invention are useful for determining CC compositions of the present invention as mad/or cell culture media. The antigens, antibodies and methods of the present invention, as CC compared to prior art, have the particular advantages of possessing CC greater affinity for FTH, and in particular advantages of possessing CC prior and further have negligible cross-reactivity with the large con-molecular forms of FTH. FTH levels are an important parameter in patients suffering from hypercalcaemia, osteoporosis and primary hypercarathyroidism. The present sequence represents a PTH antigenic copplicate representing amino acids 2-12 of human, mouse, rat, porcine, CC canine and bovine PTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                  Mus
                          (ZAHR/)
                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                      sus
                                                                                                                                                                                                                                                                                                                                                                                            Human; parathyroid hormone; antigen; hypercalcaemia; osteoporosis; primary hyperparathyroidism; mouse; rat; bovine; porcine; canine.
                                                                     05-DEC-2000; 2000US-00730174.
                                                                                                   05-DEC-2000; 2000US-00730174
                                                                                                                                   15-AUG-2002
                                                                                                                                                                                                                                          Misc-difference
                                                                                                                                                                                                                                                                                                                    Bos taurus.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Parathyroid hormone antigenic peptide 1-12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        11-FEB-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABG72608 standard; peptide; 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 11
                                                                                                                                                               US2002110871-A1
                                                                                                                                                                                                                                                                                                                                 Rattus sp
                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                      scrofa.
is familiaris
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10; Conserv
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                          ZAHRADNIK R J.
LAVIGNE J R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VSEIQFMHNLG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       N-terminus of parathyroid hormone
                                                                                                                                                                                              /label= Leu,
                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                         /label= Ser, Ala
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12
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Pred. No. 0.01
0; Mismatches
                                                                                                                                                                                              Phe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 6;
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Sequence 12 AA;

0

bovine

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CC are; (1) a method for producing antibodies useful in the determination of CPTH levels in a biological sample comprising: (a) providing at least one C first peptide antigen comprising a peptide fragment of PTH; (b) CC administering the first peptide antigen to a host animal to induce CC antibody production; (c) monitoring the antibody titre produced; (d) CC isolating antisera produced in the host animal; and (e) Selecting CC antisera from the isolated antisera produced in the host that is capable CC the method; and (3) test kits and analytical procedures used for the CC the method; and (3) test kits and analytical procedures used for the CC compositions of the present invention are useful for determining CC bioactive intact PTH levels in serum, plasma and/or cell culture media. CC The antigens, antibodies and methods of the present invention, as CC compared to prior art, have the particular advantages of possessing CC greater affinity for PTH, and in particular, are designed to have a novel CC proposition for maino acid residues extending beyond the first N-terminal CC PTH residue, and further have negligible cross-reactivity with the large connected to procedure sequence represents a PTH antigenic paptide representing amino acids 1-12 of human, mouse, rat, porcine,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New parathyroid hormone (PTH) antigenic peptide inducing the formation and isolation of antibodies having an affinity to it, useful for determining bloactive PTH levels in serum, plasma and/or cell culture
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a new antigenic peptide for inducing the formation and isolation of antibodies having an affinity to it, be formed from the N-terminus of parathyroid hormone (PTH). Also inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 2; Page 5; 11pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2003-066685/06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Also included
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             being
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RESULT 3
AAY96968
ID AAY96968
AC AAY9
XX AAY9
XX PTH,
KW PTH,
KW bone
XX bone

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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
(GARD/) GARDELLA T J.
(KRON/) KRONENBERG H M.
(POTT/) POTTS J T.
                                                                                                                                                                  31-DEC-1998;
                                                                                                                                                                                                                                                 30-DEC-1999;
                                                                                                                                                                                                                                                                                                                                     06-JUL-2000.
                                                                                                                                                                                                                                                                                                                                                                                                               WO200039278-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Parathyroid hormone N-terminal signaling domain (residues 1-11).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            31-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAY96968 standard; peptide; 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             parathyroid hormone; N-terminal; signaling domain; bone mass; reformation; resorption; remodeling; tether1; osteoporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 VSEIQFMHNLG 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first
                                                                                                                                                                      98US-0114577P
                                                                                                                                                                                                                                                     99WO-US031108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       entry)
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Pred. No. 0.014;
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Zahradnik RJ

Lavigne JR;

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AXAXAXAAXOOOOOOOOOOOO
8
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Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Compounds of the structure or formula S-(L) n-B, R_1-S-(L) n-R or S-(L) n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4)); B is a C-terminal binding portion of pTH(1-34) or a pTH-related protein (PTH-PP) (1-34); R 1 is the PTH-1 receptor signal sequence; and R is (a portion of) pTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAMP in a mammalian cell having pTH-1 receptors, or screening for a peptide or non-peptide PTH (claimed). The new compound can be administered by regular injections to treat osteoporosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.
            Claim
                                    New antigenic peptide(s) fr
generated using them, able
forms of the hormone.
                                                                                                                                                                                                                               04-APR-1996
                                                                                                                                                                                                                                                                                                             diagnosis;
                                                                                                                                                                                                                                                                                                                          Human
                                                                                                                                                                                                                                                                                                                                                 Human parathyroid hormone antigenic peptide hPTH 1-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                        28-SEP-1994;
                                                                                                                                                                                                                                                         DE4434551-A1
                                                                                                                                                                                                                                                                                                                                                                                                           AAR91644,
                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR91644 standard;
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                                                                                          WPI; 1996-180391/19
                                                                                                                                                (FORS/)
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           2;
                                                                                                                                                                                                                                                                                                                       parathyroid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10;
                                                                                                                                                FORSSMANN W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
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          Page 4; 5pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AVSEIQFMHNL 11
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                                                                                                                                                                          94DE-04434551.
                                                                                                                                                                                                    94DE-04434551
                                                                                                                                                                                                                                                                                                             hormone; hPTH; antigen; alpha-helix; hPTH 1-37.
                                                                                                                                                                                                                                                                                                                                                                                                                                   peptide; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90.3%;
            German
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Pred. No. 0.02;
                                                  human parathyroid hormone - and antibodidistinguish between active and inactive
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Query Match Best Local S Matches 8

Similarity 8; Conserv

Conservative

68.9%;

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Best Local S
Matches 8
                                                                                        The present sequence represents a parathyroid hormone (pTH) fragment, comprising amino acids 1-10. Parathyroid hormone fragments of the invention have lipolysis stimulating activity (especially when topically administered). The lipolytic activity of the peptides is enhanced when they are chemically modified to increase their lipolyhilicity. The peptides are used in cosmetic or dermatological compositions for skin care. They are especially used for slimming treatment of excessive weight in the thighs and hips, in the treatment of cellulite and for skin
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                                                                                                                                                                                                                                                                                                                                                                                                                                     New parathyroid hormone fragment peptides, in topically applied cosmetic compositions excessive weight in hips and thighs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17-JUL-1998;
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8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              parathyroid hormone (pTH)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 42; DB
Pred. No. 0.37
1; Mismatches
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0.37;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         used as lipolysis stimulants for slimming treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 10,
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RESULT 6
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RESULT 7
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                                                                                                                                                                                                                                                                                                        This invention describes a novel method for determining (M1) the content of active parathyroid hormone (A) by treating a sample with (1) antibody (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), and including the N-terminal residue and (ii) antibody (Ab2) that creognizes an epitope within the receptor-binding site of (A). The number of molecules that react with both antibodies is determined and used to calculate the content of physiologically active (A). The method is used for diagnosis of calcium-metabolism disorders, osteopathy, or hypo- or hyper-parathyroidism. The method (unlike known assays) recognizes that (C) some fragments of (A) shorter than the complete (B4 aa) peptide are active and (ii) that apparently intact peptide may be biologically contactive, and also takes into account the fact that some fragments of (A) care antiagonistic (these have the receptor-binding site but lack the N-CC terminus). It thus provides a true measure of the content of physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This core the invention appears to a peptide fragment used to illustrate the method of
                                                                                                                                                                                                Query Match
Best Local S
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibodies reactive with different epitopes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist;
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                   ABR44166;
                                                  ABR44166 standard; peptide;
                                                                                                                                                                                                                                                                Sequence 10 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 3; 10pp; German
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                                                                                                                                                                                                                                                                                                 invention
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8; Conservative
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                                                                                                                                                                                                Score 42; DB 4;
Pred. No. 0.37;
1; Mismatches
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Matches
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Chang M;
                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a fusion peptide (Tat-hPTHDP), where self cell-penetrating Tat peptide is bound to human parathyroid hormone-derived peptide (hPTHDP). The fusion peptide is useful as a component of skin slimming cosmetic composition. The fusion peptide does not cause irritation, easily and safely penetrates into integument and endothelium, does not cause skin disease and has superior lipolysis effects, and is durable. The present sequence represents a human parathyroid hormone (hPTH) fragment that can be used to construct the fusion peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel fusion peptide comprising self to human parathyroid hormone-derived slimming cosmetic composition.
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                                                                                                                                                                                                                                                                                                                                                                                               Sequence 10 AA;
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15-MAR-2002; 2002KR-00014062.
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lipolysis; human; hPTH.
                                                                                             Cyclic peptide; resin bound; parathyroid hormone; osteopathic; disea hypotensive; PTH receptor; treatment; hyper-calcemia; hypo-calcemia; osteoporosis; osteopenia; hyper-parathyroidism; hypo-parathyroidism; Cushing's syndrome; renal failure; hypertension; bone fracture repai
                                                                                                                                                                                                                                   AAY50600 standard; peptide; 11 AA
             Misc-difference
                                                                                                                                                            Resin bound
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                                    Misc-difference
                                                                         Synthetic
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8; Conserv
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                                                                                                                                                            cyclic peptide 33
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                                             Location/Qualifiers
/note= "Ser(OtBu)"
                       /note=
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                                                                                                                                                                                                                                                                                                                                                          Score 42; DB 6;
Pred. No. 0.37;
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                                                                                                                                                                                                                                                                                                                                                  Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cell-penetrating Tat peptide peptide, useful as component
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Lee Y,
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of skin
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Misc-difference

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                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel method for the preparation of peptides (II) that contain both cyclic and linear peptide fragments comprises (c sequential reaction of a resin-bound linear fragment with the cyclic fragment in N-protected form and optionally other linear fragments. The products of the invention have osteopathic and hypotensive activity. (II) think the cyclic method is particularly used to prepare cyclic peptide analogs of parathyroid hormone (PTH) or PTH-related peptides which are useful for treating diseases that respond to treatment with agents that bind to PTH creoptors (with or without activation of adenyl) cyclase activity), e.g. hyper- or hypo-calcemia, osteoporosis, osteopenia, hyper- or hypo-calcemia, osteoporosis, osteopenia, hyper- or hypo-calcemia fractures. Separate synthesis of the cyclic for promoting repair of bone fractures. Separate synthesis of the cyclic fragment allows convergent synthesis of resin-bound (II), with better cyields and higher throughput. The difficulties associated with the peptide before reaction with the resin-bound component. AAY50568-creation of the bridged fragment are confined to a small peptide which can be purified before reaction with the resin-bound component. AAY50568-creation of the bridged fragments described in the method of the
                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local S
Matches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
                                               Parathyroid hormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist;
                                                                                           Human parathyroid
                                                                                                                               03-SEP-2001
                                                                                                                                                              AAB86225;
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                              hypo-parathyroidism;
                                                                                                                                                                                            AAB86225 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (RHON ) RHONE-POULENC RORER PHARM INC
                                                                                                                                                                                                                                                                                               1 AVSEIQFMHNLG 12
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                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                         AVSEIQ-LHNLG 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              synthesis of peptides for treating e.g. bone disorders
                                                                                                                                                                                                                                                                                                                                                                                                       ĀΑ;
                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                             (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mencel
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        note=
                                                                                               hormone
                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                       68.0%;
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                              hyper-parathyroidism
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Asn(Trt)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Gln(Trt)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "His(Trt)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Glu (OtBu)
                                                                                             immunogenic peptide SEQ
                                                                                                                                                                                                                                                                                                                                                         Score 41.5;
Pred. No. 0
                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                        DB
                                                                                                                                                                                                                                                                                                                                                                       2:
                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                       Length 11;
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                                                                                                                                                                                                                                                                                                                                          Indels
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RESULT 10
AAB01862
ID AAB01
XX AAB01
XX I1-SE
XX Parat
KW Calci
KW Calci
KW Calci
KW Cone
XX Homo
OS Synth
XX WO200
XX Y 27-AE
XX 20-OC

11-SEP-2000 AAB01862;

(first entry)

PTH(1-14)/PTHrP(1-14)-derived peptide,

SEQ ID

NO: 6

AAB01862 standard; peptide; 9

27-APR-2000.

20-OCT-1999;

99WO-US024481

Synthetic Homo sapiens

WO200023594-A1

Parathyroid hormone peptide; PTH; PTH-related peptide; PTHrP; calcium homeostasis; PTH-1 receptor; PTH-2; vitamin D synthes bone synthesis; agonist; osteoporosis; non-parenteral deliver

synthesis; delivery.

밁 Ş

1 VSEIQLMHN 9 2 VSEIQFMHN 10 Similarity 8; Conserv

Matches Query Match

Conservative

88.9%;

Score 41; DB 4; L. pred. No. 1.4e+06; 0; Mismatches 1;

Length 9; Indels

0; Gaps

0

Local

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of active parathyroid hormone (A) by treating a sample with (i) antibody (C (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), and including the N-terminal residue and (ii) antibody (Ab2) that c recognizes an epitope within the receptor-binding site of (A). The number of molecules that react with both antibodies is determined and used to calculate the content of physiologically active (A). The method is used for diagnosis of calcium-metabolism disorders, osteopathy, or hypo- or c hyper-parathyroidism. The method (unlike known assays) recognizes that c (i) some fragments of (A) shorter than the complete (84 aa) peptide are active and (ii) that apparently intact peptide may be biologically c inactive, and also takes into account the fact that some fragments of (A) are antagonistic (these have the receptor-binding site but lack the N-c terminus). It thus provides a true measure of the content of physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This sequence represents a peptide fragment used to illustrate the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibod reactive with different epitopes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes a novel method for determining (M1) the content of active parathyroid hormone (A) by treating a sample with (i) antibody (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A).
Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 3; 10pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-376318/40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-DEC-1999;
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RESULT 11
AAY96966
ID AAY96
XX
AC AAY96
XX
DT 31-OC
DT 31-OC
DX
DE Parat
XX
KW PTH;
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                                                                                                                                                                                                                                                                                                                                                                                                                          CC generic peptide of the formula: X1 Val-Ser-Glu-X2-Gln-Leu-X3-His-X4-X5-CC Gly-Lys-X6 (AAB01857) where: X1 is Ser or Ala; X2 is Ile or Lys; X3 is CC Mt, Leu or Nle; X4 is Asn or Asp; X5 is Leu or Lys; X6 is His or Ser; CC provided that the peptide is not PTH-P11-14). The peptides of the CC invention also encompass fragments of peptides of the invention CC consisting of amino acids 1-9, 1-10, 1-11, 1-12 and 1-13, and N- and C- terminal derivatives. PTH is a major regulator of calcium homeostasis, CC and is necessary for the normal function of the gastrointestinal, steletal, neurological system, neuromuscular and cardiovascular systems. CC the binds to both PTH-1 receptors on osteoblasts and renal tubular cells; CC and to the recently identified PTH-2 receptor. PTH has a potent anabolic ceffect on the skeleton, and mediates calcium reabsorption, enhances composite the renal and skeletal actions of PTH, and also bind to the PTH-1 creceptor. The peptides of the receptor. They do not bind to the PTH-2 receptor. The peptides of the invention are either agonists of PTH-1 and PTH-2 receptors (AAB01858, AB01869). PTH-1/PTH-2 receptor agonists are useful for the treatment of CC conditions characterised by a decrease in bone mass, such as condical disorders that arise from excessive or altered action of the PTH-CC majorists are useful for the treating consecution of the pth-1 pth-2 receptor. Detectably labelled peptides of the invention are also consecution of pth or PTH-P which are inexpensive to a patient via conventional synthetic chemistry, and can be delivered to a patient via conventional synthetic chemistry, and can be delivered to a patient via conventional synthetic chemistry, and can be delivered to a patient via convention are excessor.
                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a novel parathyroid hormone (PTH) peptide (AAB01859) and parathyroid hormone-related peptide (PTHrP; AAB01860-biologically active derivatives thereof (AAB01857-B01858, AAB01861-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Parathyroid hormone acids that encode th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gardella TJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (KRON/)
                                    Parathyroid hormone N-terminal signaling domain (residues 1-9).
                                                                        31-OCT-2000
                                                                                                          AAY96966
                                                                                                                                             AAY96966 standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                Sequence 9
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POTTS J T.
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8
                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                       AVSEIQFMH 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The peptides of the invention are at least 85% identical
                                                                                                                                                                                                                                    AVSEIQLMH
                                                                                                                                                                                                                                                                                                                                                                                  AA
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                                                                                                                                                                                                                                                                                                                                                                                                                receptor agonists
                                                                                                                                                                                                                                                                                                           Conservative
                                                                        (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kronenberg
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   hormone; N-terminal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ne (PTH) peptides,
them, useful for (
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                                                                      entry)
                                                                                                                                                                                                                                                                                                                            63.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 73pp; English.
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                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                            Score 39; DB 3; Pred. No. 1.4e+06;
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                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      treating osteoporosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PTH related peptides and the nucleic
   signaling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Jueppner
                                                                                                                                                                                                                                                                                                           1; Indels
                                                                                                                                                                                                                                                                                                                                             Length 9;
   domain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               H
   bone mass
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            peptide
; AAB01860),
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                                                                                                                                                                                                                                                                                                         Gaps
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RESULT 12
AAR91645
ID AAR91
XX AAR91
XX AAR91
XX Humar
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC compounds of the structure or formula S-(L) n-B, R 1-S-(L) n-R or S-(L) n CC -R, are new. S is an amino terminal signaling functional domain of the grathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4)); B is a C-terminal binding CC portion of PTH(1-34) or a PTH-related protein (PTH-rP) (1-34); R 1 is the CC PTH-11 receptor signal sequence; and R is (a portion of) PTH-1 receptor Sequence. The new compounds are used for treating mammalian conditions CC characterized by decreases in bone mass, determining rates of bone conformation, bone resocrated with decreased techer1 activity, increasing cAMP in a mammalian cell having PTH-1 receptors, or screening for a peptide or con-peptide PTH (claimed). The new compound can be administered by cregular injections to treat osteoporosis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          bone
                                                                                                                                                                  Synthetic
                                                                                                                                                                                                                  Human parathyroid hormone; hPTH; antigen; alpha-helix; immunogen; diagnosis; active hPTH 1-37.
                                                                                                                                                                                                                                                                                                        Human parathyroid
                                                                                                                                                                                                                                                                                                                                                                   06-NOV-1996
                                                                                                                                                                                                                                                                                                                                                                                                                        AAR91645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 9 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 4; Page 92; 119pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          reformation; resorption; remodeling; tether1; osteoporosis.
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) POTTS J T.
) JUEPPNER H.
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8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard; peptide; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AVSEIQFMH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AVSEIGLMH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Kronenberg
                                                                                                                                                                                                                                                                                                                                                                (first entry
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                                                                                                                                                                                                                                                                                                        hormone antigenic peptide hPTH 1-9.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         φ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           63.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 39; DB 3;
Pred. No. 1.4e+06;
0; Mismatches 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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04-APR-1996

DE4434551-A1

28-SEP-1994;

94DE-04434551

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RESULT 13
AAB01863
ID AAB01
XX AAB018
AC AAB01
XX AAB01
XX Parat
KW Parat
KW Calci
KW Calci
KW Calci
XX Synth
XX 27-Ai
XX 22-O(
XX 22-O(
XX (GARI
PA (GARI
PA (GARI
PA (GUE)
XX (FOT)
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XX (FOT)
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XX (FOT)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Parathyroid hormone peptide; PTH; PTH-related peptide; calcium homeostasis; PTH-1 receptor; PTH-2; vitamin D s bone synthesis; agonist; osteoporosis; non-parenteral d
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence is a specific example of claimed immunogenic peptides having a sequence from hPTH(1-37) which includes the N-cterminal alpha-helical region and/or the non-structured region of hormone. Antibodies and their binding fragments generated by injection with the peptides are useful as diagnostic reagents for determination of biologically active hPTH(1-37)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim
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                                                                                            Parathyroid hormone (PT) acids that encode them,
                                                Disclosure; Page 26;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                                                     Gardella TJ,
                                                                                                                                                                                                                                                                                             (KRON/)
                                                                                                                                                                                                                                                                                                                                                                                             22-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PTH(1-14)/PTHrP(1-14)-derived peptide, SEQ ID NO:7.
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                                                                                                                                                                         2000-339693/29
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                                                                                                                                                                                                                                                                                           GARDELLA T J.
KRONENBERG H M.
POTTS J T.
                                                                                                                                                                                                                                                                    JUEPPNER H.
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                                              73pp; English
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                                                                                               H) peptides,
useful for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 36; DB 2;
Pred. No. 1.4e+06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         human parathyroid hormone - and antibodies distinguish between active and inactive
                                                                                                                                                                                                                     Potts
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                                                                                                                                                                                                                   JT,
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                                                                                                                      PTH related peptides and the nucleic
                                                                                                                                                                                                                   Jueppner
                                                                                               osteoporosis
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delivery.
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novel parathyroid

hormone

(PTH) peptide

Gardella

Kronenberg

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Potts ŢŢ,

Jueppner

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JUEPPNER H. ΤJ,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Best Local
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                                                                                                                                      31-DEC-1998;
                                                                                                                                                                         30-DEC-1999;
                                                                                                                                                                                                                06-JUL-2000.
                                                                                                                                                                                                                                                        WO200039278-A2.
                                                                                                                                                                                                                                                                                                                                                                                          Parathyroid hormone N-terminal signaling domain.
                                                                                                                                                                                                                                                                                                                                                                                                                                 31-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAY96981;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 9 AA;
                                                       (GARD/) GARDELLA T J. (KRON/) KRONENBERG H (POTT/) POTTS J T.
                                                                                                                                                                                                                                                                                           sapiens
                                                                                                                                                                                                                                                                                                                                parathyroid hormone; N-terminal; signaling domain; bone mass; reformation; resorption; remodeling; tetherl; osteoporosis.
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                                                                                                                                      98US-0114577P
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77.8%;
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1.4e+06;
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RESULT 15
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Best Local S
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Compounds of the structure or formula S-(L)_n-B, R_1-S-(L)_n-R or S-(L)_n-R, are new. S is an amino terminal signaling functional domain of parathyroid hormone (PTH); L is a linker molecule present n times (where n is a positive integer 1-10 (preferably 4)); B is a C-terminal binding portion of PTH(1-34) or a PTH-related protein (PTH-TP) (1-34); R_1 is the PTH-1 receptor signal sequence; and R is (a portion of) PTH-1 receptor sequence. The new compounds are used for treating mammalian conditions characterized by decreases in bone mass, determining rates of bone reformation, bone resorption and/or bone remodeling, treating diseases and disorders associated with decreased tether1 activity, increasing cAMP in a mammalian cell having PTH-1 receptors, or screening for a peptide or non-peptide PTH (claimed). The new compound can be administered by inhalation unlike the large native PTH or PTHrP which avoids the need for regular injections to treat osteoporosis
                                                                             Determining the content of physiologically active parathyroid hormone, useful in diagnosis of calcium-metabolism disorders, using two antibodies reactive with different epitopes.
                                                                                                                                              WPI;
                                                                                                                                                                                                                                                                                                                                                                                                 Parathyroid bormone; hPTH; immunogenic peptide; human; epitope; diagnosis; calcium-metabolism disorder; osteopathy; antagonist; hypo-parathyroidism; hyper-parathyroidism.
                                                                                                                                                                             Armbruster FP;
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                                                                                                                                                                                                                                                                         17-DEC-1999;
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                                                                                                                                                                                                                                                                                                                                                                      Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human parathyroid hormone immunogenic peptide SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     03-SEP-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New compound comprising an amino terminal signaling functional domain linked to a carboxy-terminal binding portion of parathyroid hormone for treating mammalian conditions characterized by decreases in bone mass.
                                                                                                                                                                                                           (IMMU-)
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7; Conservative
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Disclosure; Page 3; 10pp; German

parathyroid hormone (A)

method for determining (M1) the content) by treating a sample with (i) antibody

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                                                                                                                                                                                                                                 CC (Ab1) that recognizes the epitope formed by amino acids (aa) 1-3 of (A), CC and including the N-terminal residue and (ii) annibody (Ab2) that CC recognizes an epitope within the receptor-binding site of (A). The number of molecules that react with both antibodies is determined and used to CC calculate the content of physiologically active (A). The method is used CC for diagnosis of calcium-metabolism disorders, osteopathy, or hypo- or CC hyper-parathyroidism. The method (unlike known assays) recognizes that CC active and (ii) that apparently intact peptide may be biologically CC inactive, and also takes into account the fact that some fragments of (A) in the receptor-binding site but lack the N-CC terminus). It thus provides a true measure of the content of physiologically active (A); contrast methods that measure intact peptide and its 1-37 fragment which may produce falsely high values. This CC sequence represents a peptide fragment used to illustrate the method of the invertice.
                                                                               Query Match
Best Local Similarity
Matches 7; Conserv
                                                                                                                                                                           Sequence 9 AA;
                                                                                                                                                                                                                       invention
                                          AVSEIQFMH 9
SVSEIQLMH
                                                                                    Conservative
ω
                                                                                                         59.0%;
77.8%;
                                                                                      1; Mismatches
                                                                                                         Score 36; DB 4;
Pred. No. 1.4e+06;
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Search Job tim time completed: May 18, me : 45.75 secs 2004, 10:01:54

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Result
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Maximum DB
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           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score
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seq length: 12
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Match
     US-09-730-174A-6
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100.425 Million cell updates/sec
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Gapop 10.0 , Gapext
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: /cgn2_6/ptcdata/1/pubpaa/US08_PUBCOMB.pep:*
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US-09-730-174A-3
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US-09-730-174A-1
US-09-730-174A-1
US-09-730-174A-1
US-10-168-185-7
US-10-168-185-7
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US-10-168-185-7
US-10-168-185-8
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US-09-730-174A-6
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-10-147-140-	-10-043-487-5	-10-168-185-	0-168-185-	-10-319-130-1	0-153-334-	S-10-033-741-6	9-910-059	-09-834-765-7	-09-834-765-7	-09-834-765-5	9-834-765-4	-09-791-551-1	-10-430-176-	-10-366-709-	-10-137-867-5	-10-230-880-11	-10-283-349-4	-10-160-232-1	-10-160-232-	-10-195-752-1	-10-051-852-1	-10-239-656-1	798-689-	9-09-563-222-	-965-099-1	9-865-198-	-976-787-	10-192-6	-10-366-709-
equence 26	equence 508, Ap	equence 4, Appl	equence 5, Appl	equence 16, Ap	equence 14, App	ence 61, Āp	equence 26, App	e 78, App	equence 73, App	equence 55, App	equence 45, App	e 13, App	equence 12,	equence 3,	equence 526	equence 115	equence 49,	equence 18,	equence 12,	equence 108	equence 12,	equence 19,	equence 4	equence 41,	equence 12,	quence 4, A	equence 4, A	O, Ap	equence 2, A

ALIGNMENTS

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; OTHER INFORMATION: Induces Formation of Antibodies and Isolates US-09-730-174A-6
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; Patent No. US20020110871A1
; GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
; TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having; TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid FILE REFERENCE: IMUNE-001A
Sequence 4, Application US/09730174A
Patent No. US20020110871A1
GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having
TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid H
FILE REFERENCE: IMUNE-001A
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CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 6
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                      1 AVSEIQFMHNLG 12
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RESULT 4
US-09-730-174A-5
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Best Local S
Matches 11
                                                                                                                                                              TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Hc FILE REFERENCE: INUNE-001A CURRENT APPLICATION NUMBER: US/09/730,174A CURRENT FILING DATE: 2000-12-05 NUMBER OF SEQ ID NOS: 12
                                                                                                                                                                                                                                                                             Sequence 5, Application US/09730174A PATENT NO. US20020110871A1 GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
     Matches
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CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 4
LENGTH: 12
TYPE: PRT
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having
TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid F
FILE REFERENCE: IMUNE-001A
FILE REFERENCE: IMUNE-001A
CURRENT FAPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
                                                                   FEATURE:
OTHER INFORMATION: Induces Formation
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TYPE: PRT
ORGANISM: Artificial
                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                ENGTH: 12
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91.7%;
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Score 55; DB 9;
Pred. No. 0.0015;
0; Mismatches
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Pred. No. 0.0006;
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Pred. No. 0.00043;
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Matches 10
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Best Local 9
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; Sequence 1, Application US/10168185
; Publication No. US20030175802A1
; GENERAL INFORMATION:
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APPLICANT: Zahradnik, R.J.
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Ho: FILE REFERENCE: IMUNE-001A
CURRENT APPLICATION NUMBER: US/09/730,174A
CURRENT FILING DATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3, Application US/09730174A
Patent No. US20020110871A1
GENERAL INFORMATION:
APPLICANT: Zahradnik, R.J.
APPLICANT: Zahradnik, R.J.
TITLE OF INVENTION: Antibodies and Peptide Antigens for Producing Antibodies having a TITLE OF INVENTION: Selective Binding Specificity to Bioactive Intact Parathyroid Ho: FILE REFERENCE: IMUNE-001A
CURRENT EPLICATION NUMBER: US/09/730,174A
CURRENT FILING BATE: 2000-12-05
NUMBER OF SEQ ID NOS: 12
SEQ ID NO 3
LENGTH: 12
TYPE: PRT
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                          APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrici
APPLICANT: Roth, Heinz-Jurgen
APPLICANT: Roth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 11
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Induces Formation of Antibodies and Isolates said Antibodies
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                        Missbichler, Albert
Schmidt-Gayk, Heinrich
Roth, Heinz-Jurgen
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Pred. No. 0.0078;
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Pred. No. 0.0055;
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PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER OF SEO ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1
SEQ ID NO 1
LENGTH: 10
TYPE: PRT
ORGANISM: Homo sapiens
US-10-168-185-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR PILING DATE: 1999-12-17
PRIOR FILING DATE: 1999-12-17
NUMBER: OF SEQ ID NOS: 11
SOFTWARE: FRSESEQ for Windows Version 4.0
SEQ ID NO 7
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US-10-192-673-6
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Best Local S
Matches 8
                                                                                                          Sequence 6, Application US/10192673
Publication No. US20030166838A1
GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 7, Application US/10168185
Publication No. US20030175802A1
GENERAL INFORMATION:
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Best Local 9
APPLICANT: Juppner, Harald TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid TITLE OF INVENTION: Hormone-Related Peptide (PTHrp) FILE REFERENCE: 0609.4570002
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APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Juzgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
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CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
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TYPE: PRT
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Local Similarity 80.0%;
les 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 41; DB 14
Pred. No. le+06;
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Pred. No. 0.33;
1; Mismatches
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APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roch, Heinz-Curgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: PGT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 199912-17
NUMBER OF SEQ ID NOS: 11
SECTIMARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 6
                                                                                                                                                                                                                       US-10-192-673-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-10-168-185-8
                                                                                                                                                  Sequence 7, Application US/10192673
Publication No. US20030166838A1
GENERAL INFORMATION:
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Matches
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Publication No. US20030175802A1
GENERAL INFORMATION:
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APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTH)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Homo sapiens
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8; Conservative
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Pred. No. 1e+0
0; Mismatches
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Pred. No. 1e+06;
0; Mismatches
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le+06;
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APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Reth, Heinz-Jurgen
TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
TILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: DCTEP00/12911
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-112-17
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Best Local Similarity
7; Conserve
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                                                                                                                                                                                                                         US-10-192-673-8
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                                            Sequence 8, Application US/10192673
Publication No. US20030166838A1
GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
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Best Local Similarity
Thes 7; Conserv
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION
APPLICANT: Juppner, Harald TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/192,673
CURRENT FILLNG DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR FILING DATE: 1998-10-22
PRIOR FILING DATE: 1998-110-22
PRIOR FILING DATE: 1998-110-22
PRIOR FILING DATE: 1998-110-22
PRIOR FILING DATE: 1998-110-22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
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77.8%;
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77.8%;
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Pred. No.
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RESULT 15 US-10-232-187-8

Sequence 8, Application US/10232187 Publication No. US20030092091A1 GENERAL INFORMATION:

APPLICANT: Abrahamson, APPLICANT: Bochner, B

Julie A.

Bochner, Bruce Erickson-Miller, Connie L. Kikly, Kristine K. Schleimer, Robert

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CURRENT FILING DATE: 2002-07-11
PRIOR APPLICATION NUMBER: U.S. 09/421,379
PRIOR FILING DATE: 1999-10-20
PRIOR APPLICATION NUMBER: U.S. 60/105,530
PRIOR FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                    US-10-168-185-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 3, Application US/10168185 Publication No. US20030175802A1 GENERAL INFORMATION:
                                                                                Best Local Similarity
Matches 6; Conserv
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                                                                                                                          Query Match
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Best Local
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TITLE OF INVENTION: Method for Determining Parathormone
TITLE OF INVENTION: Activity in a Human Sample
FILE REFERENCE: HLZ-004US
CURRENT APPLICATION NUMBER: US/10/168,185
CURRENT FILING DATE: 2002-66-17
CURRENT FILING DATE: 2002-66-17
                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 11
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: PCT/EP00/12911
PRIOR FILING DATE: 2000-12-18
PRIOR APPLICATION NUMBER: DE 19961350
PRIOR FILING DATE: 1999-12-17
NUMBER FILING DATE: 1999-12-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Armbruster, Franz Paul
APPLICANT: Missbichler, Albert
APPLICANT: Schmidt-Gayk, Heinrich
APPLICANT: Roth, Heinz-Jurgen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Hormone-Related Peptide (FTHrP) FILE REFERENCE: 0609.4570002
                                                                                                                                                                                TYPE: PRT ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 9
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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                                        1 AVSEIGFM 8
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SVSEIOLM
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75.0%;
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Pred. No. 1e+06;
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APPLICANT: Nulku, Turkan E.

FILE OF INVENTION Sialoadhesin Factor-2 Antibodies

FILE REFERENCE: GH50042-1

CURRENT APPLICATION NUMBER: US/10/232,187

CURRENT FILING DATE: 2002-08-29

PRIOR APPLICATION NUMBER: CFUSOL/07193

PRIOR APPLICATION NUMBER: CFUSOL/07193

PRIOR FILING DATE: 2001-03-05

PRIOR FILING DATE: 2001-03-05

PRIOR FILING DATE: 2001-03-05

PRIOR FILING DATE: 2002-07-10

PRIOR FILING DATE: 2001-03-30

PRIOR FILING DA
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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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Match
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48.589 Million cell updates/sec
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Copyright (c) 1993 - 2004 Compugen Ltd.
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US-09-421-379-6
US-09-421-379-7
US-09-421-379-7
546097-3
US-09-421-379-8
US-09-421-379-8
US-08-417-547A-3
US-08-417-547A-3
US-08-418-7528-42
US-08-4813-528-42
US-08-4813-588-9
US-08-4813-588-9
US-08-4813-588-9
US-08-4813-588-108
US-09-344-050-12
US-09-345-050-12
US-09-345-050-13
US-08-817-547A-16
US-08-817-547A-16
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Sequence 31, Appl
Sequence 6, Appli
Sequence 7, Appli
Sequence 7, Appli
Sequence 8, Appli
Sequence 8, Appli
Sequence 42, Appli
Sequence 42, Appli
Sequence 42, Appli
Sequence 42, Appli
Sequence 12, Appli
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                         US-09-442-989-31
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                                                                                                                     FEATURE:

NAME/KEY: PEPTIDE

NAME/KEY: (10)
                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
NAME/KEY: PEPTIDE
LOCATION: (3)
OTHER INFORMATION: 8
                                      LOCATION: (8)
OTHER INFORMATION:
                                                                   REATURE: MOD_RES
                                                                                                        LOCATION: (10)
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                   FEATURE:
NAME/KEY: PEPTIDE
                                                                                                                                                                              OTHER INFORMATION:
                                                                                                                                                                                                                                                  OTHER
                                                                                                                                                                                                                                                                                 NAME/KEY: PEPTIDE
                                                                                                                                                                                              LOCATION:
                                                                                                                                                                                                               NAME/KEY: PEPTIDE
                                                                                                                                                                                                                                                INFORMATION:
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ALIGNMENTS

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Sequence 31, Application US/09442989

Patent No. 665993

APPLICANT: Mencel, James W.
APPLICANT: Mencel, James J.
A
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Sequence 6, Application US/09421379
Patent No. 6495662
GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
                                                                                                                                                                                                               RESULT 2
US-08-817-547A-1
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                                                                                                         RESULT 3
US-09-421-379-6
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                                                                                                                                                                                                                                                                                                                                         TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/817,547A
FILING DATE: US/08/817,547A
FILING DATE: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: FROST, ROGER T.
REGISTRATION NUMBER: 07826-0007
TELEPAN: 404-818-3700
TELEPAN: 404-818-3700
TELEPAN: 404-818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches
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Best Local Similarity
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APPLICANT: I
APPLICANT: N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
NUTHER OF SEQUENCES: 36
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                      ANTI-SENSE: no
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: UZIP: 30303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Atlanta
STATE: Georgia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE:
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5. 6030790
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191 Peachtree Street, 37th Floor
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Hock, Dieter
                                                                                                                                                                                                                              Conservative
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                                                                                                                                                                                                                                                                                                                      no
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                                                                                                                                                                                                                                          68.9%;
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Pred: No.
                                                                                                                                                                                                                                             Pred. No. 0.
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RESULT 4
US-08-817-547A-2
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Best Local Similarity
Matches 8; Conserv:
MOLECULE TYPE: po

HYPOTHETICAL: no

ANTI-SENSE: no

US-08-817-547A-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PATENT NO. BUCKLEY
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Hock, Dieter
APPLICANT: Magerlein, Markus
APPLICANT: Magerlein, Markus
TITLE OF INVENTION: Peptides from the hPTH Sequence
TITLE OF SEQUENCES: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid TITLE OF INVENTION: Hormone Related Peptide (PTHrP) FILE REFERENCE: 0609.4570001

CURRENT APPLICATION NUMBER: US/09/421,379

CURRENT FILING DATE: 1999-10-20

EARLIER APPLICATION NUMBER: US. 60/105,530

EARLIER FILING DATE: 1998-10-22

RUMBER OF SEQ ID NOS: 13
                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/817,547A
FILING DATE:
PRIOR APPLICATION DATA:
PRICATION NUMBER: PCT/EP95/03757
APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 22,176
REGISTRATION NUMBER: 07826-0007
TELECOMMUNICATION INFORMATION:
                                                                                                                                                   TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 191 Peachtree Street, 37th Floor
CITY: Atlanta
STATE: Georgia
COUNTRY: USA
                                                                                                                                                                                                                TELEPHONE:
                                                                               TOPOLOGY:
                                                                                               STRANDEDNESS:
                                                                                                                                     ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30303
                                                                                                                  amino acid
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                                                                                                                                     9 amino acids
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                                                                               unknown
                                                                                                                                                                                                                  404-818-3700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jones & Askew, LLP
                                                        peptide
                                                                                               unknown
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88.9%;
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Pred. No. 3e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 9;
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GENERAL INFORMATION:
APPLICANT: Gardella, Thomas J.
APPLICANT: Gardella, Thomas J.
APPLICANT: Woronnberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
ITILE OF INVENTION: Bloactive Peptides and Peptide Derivatives of
ITILE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
ITILE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
                                                                                                                                                                                                                           PATENT NO. 5460978

; PATENT NO. 5460978
; APPLICANT: MARTIN, THOMAS J.; MOSELEY, JAI; BRUCE E.; METTEMALL, RICHARD E.H.
; BRUCE E.; METTEMALL, RICHARD E.H.
; PATENT OF INVENTION: PROTEIN ACTIVE IN HUI; HYPERCALCEMIA OF MALICIANACY-PTHRE
NUMBER OF SEQUENCES: 4
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/715,280
APPLICATION NUMBER: 199,235
FILING DATE: 14 JUN-1991
; PRIOR APPLICATION NUMBER: 199,235
FILING DATE: 09 MAY-1988
APPLICATION NUMBER:
                                        Query Match
Best Local Similarity
Thes 7; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-09-421-379-7
                                                                                                                                                                                     SEQ ID NO:3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn Ver. 2.0 SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
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                                                                                                                                                                   LENGTH: 11
                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 AVSEIQFMH 9
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AVSEIQFMHN 10
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                                                              Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                59.0%;
77.8%;
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77.8%;
                                                                                  50.8%;
70.0%;

    Mismatches

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Pred. No. 3e+05;
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                                                                                  Score 31; DB 6; Pred. No. 8.5;
                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                        HUMORAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Indels
                                                                                                    Length 11;
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                                                              Gaps
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APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Potts, John T.
APPLICANT: Potts, John T.
APPLICANT: Juppner, Harald
TITLE OF INVENTION: Bioactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: U.S. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
SOFTWAKE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Description of Artificial Sequence: synthetic; OTHER INFORMATION: peptide US-09-421-379-8
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US-08-140-137A-42
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US-09-421-379-8
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Best Local
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APPLICANT: TUDWANEN, ELAINE
APPLICANT: MASURE, H. R.

TITLE OF INVENTION: ANALOGS OF ENDOTHELIAL LEUKOCYTE
TITLE OF INVENTION: ADHESION MOLECULE (ELAM)
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: Klauber & Jackson
                                                                                                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IEM PC compatible

COMPUTER: IEM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, V.

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/140,137A

FILING DATE: 27-MAY-1994

CLASSIFICATION: 424

ATTORNEY/AGENT INFORMATION:

NAME: Tackor Ecc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
                    NAME: Jackson Esq., David A. REGISTRATION NUMBER: 26,742 REFERENCE/DOCKET NUMBER: 600 TELECOMMUNICATION INFORMATION: TELEPHONE: 201 487-5800
                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
CITY: F
                                                                                                                                                                                                                                                                                                                                        07601
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Hackensack 
: New Jersey
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                                                                600-1-096
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Pred. No. 3e+05;
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                                                                                                                                                                                                                                   Version #1.25
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RESULT 6 5460978-3

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Matches

5460978-3

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TOPOLOGY: linear

MOLECULE TYPE: peptide

DESCRIPTION: CaRD peptide sequence
US-08-140-137A-42
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US-08-817-547A-3
                                                                                                                                                                                                                              MOLECULE TYPE: per HYPOTHETICAL: no HYPO
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Best Local Similarity 62.5%;
Matches 5; Conservative
                                                                     Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/817,547A
FILING DATE:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/03757
FILING DATE: 29 SEPT 1994
ATTORNEY/AGENT INFORMATION:
NAME: PROST, ROGER T.
REGISTRATION NUMBER: 22,176
REFERENCE/DOCKET NUMBER: 07826-0007
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-18-3700
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APPLICANT: Aderma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 404-818-3799
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
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SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
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APPLICANT: Magerlein, M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: Peptides from the hPTH Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Jones & Askew, LLP
STREET: 191 Peachtree Street, 37th Floor
CITY: Atlanta
STATE: Georgia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
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5. 6030790
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1 AVSEIGEM 8
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GY: linear
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                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                          unknown
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                                                                                                               45.9%;
75.0%;
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Pred. No.
                                                                          Score 28; DB 3; Length 8; Pred. No. 3e+05; 1; Mismatches 1; Indels
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US-08-482-228-42
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No. 6017719
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Guthrie, Janice
REGISTRATION NUMBER: 35,170
REFERSNCE/DOCKET NUMBER: IT--
TELECOMMUNICATION INFORMATION:
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COMPUTER READABLE FORM:
COMPUTER: IEM PC compatible
COMPUTER: EM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
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tent No. 5966/J.
TRENGTION:
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APPLICANT:
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APPLICANT:
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TITLE OF INVENTION: POSITIVE AND POSITIVE/NEGATIVE
TITLE OF INVENTION: SELECTION MEDIATED BY PEPTIDE F
NUMBER OF SEQUENCES: 215
                                                                                                           APPLICANT: Al-Abdaly, Fahad A.
APPLICANT: Guillermo, Roy
APPLICANT: Helgerson, Sam L.
APPLICANT: Deans, Robert J.
TITLE OF INVENTION: POSITIVE AND POSITIVE/NEGATIVE CELL
TITLE OF INVENTION: SELECTION MEDIATED BY PEPTIDE RELEASE
NUMBER OF SEQUENCES: 215
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                                                                              CORRESPONDENCE ADDRESS:
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TOPOLOGY:
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ADDRESSEE: Janice Guthrie, Ph.D. STREET: P.O. Box 15210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
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5; Conserv
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GY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12 amino acids
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Deans, Robert J.
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                                                                                                                                                                                                                                                                                                                                                                                                 Kobori,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tseng-Law, Janet
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (714) 440-5353
                                                                                                                                                                                                                                                                                                                                                                                             Joan A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               44.3%; Score 27; DB 2; 55.6%; Pred. No. 54; tive 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IT-4630CIP3
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STATE: C

California

USA

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APPLICANT: Gardella, Thomas J.
APPLICANT: Kronenberg, Henry
APPLICANT: Potts, John T.
APPLICANT: Dotts, John T.
APPLICANT: Dotts, John T.
APPLICANT: Dotts, John T.
APPLICANT: Doppner, Harald
TITLE OF INVENTION: Bicactive Peptides and Peptide Derivatives of
TITLE OF INVENTION: Parathyroid Hormone (PTH) and Parathyroid
TITLE OF INVENTION: Hormone-Related Peptide (PTHrP)
FILE REFERENCE: 0609.4570001
CURRENT APPLICATION NUMBER: US/09/421,379
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US. 60/105,530
EARLIER FILING DATE: 1998-10-22
NUMBER OF SEQ ID NOS: 13
                                                                                                                                                            ; ORGANISM: Artificial Sequence ; FEATURE: OTHER INFORMATION: Description of Artificial Sequence: synthetic ; OTHER INFORMATION: peptide US-09-421-379-10
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                                                             Query Match
Best Local Similarity
Matches 5; Conserv
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Best Local Similarity 55.6%;
Matches 5; Conservative
                                                                                                                                                                                                                                                                                   SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
LENGTH: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 10, Applicated No. 6495662
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

COMPUTER READABLE Floppy disk

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/482,528

FILING DATE: 07-JUN-1995

CLASSIFICATION: 435

CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (714) 553-1952
INFORMATION FOR SEQ ID NO: 42:
SEQUENCE CHARACTERISTICS:
LENGTH: 12 amino acids
                                                                                                                                                                                                                                                                      TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION: TELEPHONE: (714) 440-5353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Guthrie, Janice
REGISTRATION NUMBER: 35,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AVSEIQFMH 9
                                     AVSEIQFMH 9
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SVSEHQLLH 9
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                                                                                 Conservative
                                                                                                   42.6%;
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                                                                                                   Score 26; DB 4;
Pred. No. 3e+05;
                                                                                 Mismatches
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                                                                                 Indels
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STREET: CITY: F

ARLINGTON

E: VIRGINIA TRY: U.S.A. 22201-4714 ADDRESSEE:

1100 NORTH GLEBE ROAD

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US-08-116-778E-9
                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local S
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APPLICANT: NAKAMUR
APPLICANT: KOIKE,
APPLICANT: SHITARA
APPLICANT: HANAI,
                                                                                                                                                                                 Sequence 9, Application US/08438562
Patent No. 5874255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 9, Application US/08116778E Patent No. 5830470
                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: FLOSDY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/116,778E
FILING DATE: 07-SEP-93
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: WILSON, MARY J.
REGERENCE/DOCKET NUMBER: 249-59
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (703)816-4000
TELEFAX: (703)816-4100
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: peptide
APPLICANT: HASBGAWA, MANORU
TITLE OF INVENTION: HUMANIZED ANTIBODIES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: NIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: HASEGAWA, MAMORU
TITLE OF INVENTION: HUMANIZED ANTIBODIES
NUMBER OF SEQUENCES: 49
                                                                                                            APPLICANT: NAKAMURA, KAZUYASU
APPLICANT: KOIKE, MASAMICHI
APPLICANT: SHITARA, KENYA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: ARLINGTON STATE: VIRGINIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RY: U.S.A.
22201-4714
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAKAMURA, KAZUYASU
KOIKE, MASAMICHI
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                                                                          SHITARA, KENYA
HANAI, NOBUO
KUWANA, YOSHIHISA
                                                                                                                                                                                                                                                                                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                 41.0%; Score 25; DB 2; Le 44.4%; Pred. No. 1.1e+02; ative 2; Mismatches 3;
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JENGTH: 10 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-438-562-9
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               APPLICATION NUMBER: US/08/483,528B
FILING DATE: 07-JUN-95
CLASSIFICATION: 536
TELEPHONE: (703)816-4000
TELEPHONE: (703)816-4100
INFORMATION FOR SEQ ID NO: 97:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 41.0%;
Best Local Similarity 44.4%;
Matches 4; Conservative
                                                                                                                                                                                                 ADDRESSE: NIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD
CITY: ARLINGTON
STATE: VIRGINIA
COUNTRY: U.S.A.
ZIP: 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPARTING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNIMPED. 115/08/483 5288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 97, Applicat
Patent No. 5939532
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/438,562
FILING DATE: 10-MAY-95
CLASSIFICATION: 424
PRIOR APPLICATION DATA: 108/116,778
APPLICATION NUMBER: 08/116,778
FILING DATE: 07-SEP-93
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: WILSON, MARY J.
RADISTRATION NUMBER: 32,955
REFERENCE/DOCKET NUMBER: 249-76
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: HASEGAWA, MITTLE OF INVENTION: HUINUMBER OF SEQUENCES: 1:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy
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amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       T: KUWANÀ, YOSHIHISA
T: HASEGAWA, MAMORU
INVENTION: HUMANIZED ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Application US/08483528B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAKAMURA, KAZUYASU
KOIKE, MASAMICHI
SHITARA, KENYA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HANAI, NOBUO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          103
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Pred. No. 1.1e+02;
2; Mismatches 3
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; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-483-528B-97

Query Match
Best Local Similarity 41.0%; Score 25; DB 2; Length 10;
Best Local Similarity 44.4%; Pred. No. 1.1e+02;
Matches 4; Conservative 2; Mismatches 3; Indels 0; Gaps 0;

OY 1 AVSIOFMH 9
Db 2 ASSVSYMH 10

Search completed: May 18, 2004, 10:06:44

Job time : 12.75 secs
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